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Minimum
Maximum
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Perfect score:
                                                        Database
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Maximum Match 100%
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                                                                                                                                                                                                                                                                                    Sequence:
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DB seq length: 2000000000
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                                                                                                                                                                                                                                                                                                 US-10-019-439-1
57
4 2 2 1
                                                                                                                                                                                                                                                                                                                                        June 27, 2003, 17:15:16; Search time 14.3 Seconds (without alignments) 73.950 Million cell updates/sec
                                                                                                                                                                                                             283224 segs, 96134422 residues
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Gapop 10.0 , Gapext 0.5
          PIR_73:*
1: pir1:*
2: pir2:*
3: pir3:*
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Copyright
                                                                                                                                                                                                                                                                                                                                                                                                                                           GenCore version 5.1.6 (c) 1993 - 2003 Compugen Ltd.
                                                                                    summaries
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14	13	12	11	10	9	80	7	σ	5	4	ω	2	μ.	No.	Result
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alpha-glucosidase	alpha-glucosidase	hypothetical prote	probable propionyl	heparin cofactor I	nicotinate phospho	aminomethyltransfe	syntaxin 8 related	myb-related transc		hypothetical prote		hypothetical prote	histidinol dehydro	senescence marker		-	DNA primase [impor	Ξ		prot		cifier	fibrinogen alpha c		alpha	alpha	fibrinogen alpha c	fibrinogen alpha c	Description	

A.Status: translated from GB/EMBL/DDBJ
A;Molecule type: mRNA
A;Residues: 110-156 <RES>
A;Cross-references: GB:K02272; NID:g182427; PIDN:AAA52428.1; PID:g182428
A;Cross-references: GB:K02272; NID:g182427; PIDN:AAA52428.1; PID:g182428
A;Cross-references: GB:K02272; NID:g182427; PIDN:AAA52428.1; PID:g182428
A;Fu, Y.; Weissbach, L.; Plant, P.W.; Oddoux, C.; Cao, Y.; Liang, T.J.; Roy, S.N.; Redm
Biochemistry 31, 11968-11972, 1992
A;Title: Carboxy-terminal-extended variant of the human fibrinogen alpha subunit: a nov
A;Reference number: A44234; MUID:93090725; PMID:1457396
A;Accession: A44234; MUID:93090725; PMID:1457396

A;Molecule type: mRNA A;Residues: 1-51 <FUl> A;Cross-references: GB:M64982; NID:g458553; PIDN:AAA17055.1; PID:g458554 A;Note: sequence extracted from NCBI backbone (NCBIN:119912, NCBIN:119914,

NCBIP:119918

A;Accession: C44234

33	υ W	33	33	33	33	ω	υ U	33	33	ω ω	34	34	34	34	34
	57.9		57.9	57.9	57.9	57.9	57.9	57.9	57.9	57.9		59.6			59.6
589	589	589	587	533	478	426	352	331	263	209	3472	1539	1004	950	584
2	2	N	N	N	N	N	↦	N	N	N	N	2	N	2	N
866856	S59370	850355	D84426	H86282	E86314	B75434	AJKXQ	B82341	AH0644	AC0273	T31308	T30037	851133	B87611	S64627
probable membrane	alpha-glucosidase	alpha-glucosidase	hypothetical prote	protein F10B6.34 [F2H15.15 protein -	UDP-N-acetylglucos	glutamate-ammonia	NifR3/Smm1 family	probable exported	probable exported	hypothetical 367K	hypothetical prote	transposase Tn4652	TonB-dependent rec	alpha-glucosidase

ALIGNMENTS

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Annu. Rev. Biochem. 53, 195-229, 1984
A;Title: Fibrinogen and fibrin.
A;Reference number: A90041; MUID:84305751;
                         A;Title: Cross-linki
A;Reference number:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Title: Phosphorylation of fibrinogen by casein kinase A;Reference number: A90116; MUID:84104274; PMID:6318767 A;Contents: annotation; phosphorylation A;Note: about one-third of alpha chain molecules in block phosphorylation alpha chain molecules in block phosphorylation block phosphorylation alpha chain molecules in block phosphorylation between the phosphorylation alpha chain molecules in block phosphorylation between the phosphorylation p
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ann. N. Y. Acad. Sci. 400, Co. Fibrinogen.
A;Title: Covalent structure of fibrinogen.
A;Reference number: A90037; MUID:83254370; PMID:
A;Contents: annotation; review, disulfide bonds
A;Contents: annotation; review, Misulfide bonds
A;Contents: annotation; review, disulfide bonds
A;Contents: annotation; review, disulfi
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J. Biol. Chem. 253, 2184-2195, 1978
A;Title: Localization of the alpha-chain cross-link acceptor sites of human f. A;Reference number: A92225; MUID:78130085; PMID:632262
A;Contents: annotation; cross-linking acceptor sites
R;Cottrell, B.A.; Strong, D.D.; Watt, K.W.K.; Doolittle, R.F.
Biochemistry 18, 5405-5410, 1979
Biochemistry 18, 5405-5410, 1979
A;Title: Amino acid sequence studies on the alpha chain of human fibrinogen.
A;Reference number: A90432; MUID:80088230; PMID:518845
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      R;Doolittle,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R;Itarte, E.; Plana, M.; Guasch, M.D.; Martos, C. Biochem. Biophys. Res. Commun. 117, 631-636, 1983
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Contents: annotation; cross-linking acceptor sites R;Henschen, A.; Lottspeich, F.; Kehl, M.; Southan, C. Ann. N. Y. Acad. Sci. 408, 28-43, 1983
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Molecule type: protein
A;Residues: 20-40 <DEW>
R;Retzlos, A.D.; Markland Jr., F.S.
Thromb. Res. 52, 541-552, 1988
A;Title: A direct-acting fibrinolytic enzyme from the venom of Agkistrodon contortrix A;Reference number: A60905; MUID:89162316; PMID:3232124
                                                                                                                                                                                                                 ;Kimura,
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A;Reference number: S19297; MUID:9214382
A;Accession: S19297
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A;Residues: 20-65,'T', 67-629 <BLO>
R;Dewey, R.S.; Liesch, J.M.; Williams, H.R.; Sugg,
Biochem. J. 281, 519-524, 1992
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A;Title: Disulfide bridges in NH-2-terminal part of human A;Reference number: A94309; MUID:76225080; PMID:936108
A;Contents: variant, and disulfide bonds
A;Accession: B94309
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R;Blomback, B.; Hessel, B.; Hogg, D.
Thromb. Res. 8, 639-658, 1976
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A;Residues: 605-644 <FUZ>
A;Cross-references: GB:M64982; NID:g458553; PIDN:AAA17055.1; PID:g458554
A;Cross-references: GB:M64982; NID:g458553; PIDN:AAA17055.1; PID:g458554
A;Note: sequence extracted from NCBI backbone (NCBIP:119920)
R;Henschen, A.; Lottspeich, F.; Southan, C.; Topfer-Petersen, E.
in Protides of the Biological Fluids, Proc. 28th Collog., Peeters, H., e
a. Title: Human fibrinogen: sequence, sulfur bridges, glycosylation and s
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A;Title: Amino acid sequences studies on the alpha chain of human A;Reference number: A90433; MUID:80088231; PMID:518846

A;Contents: disulfide bonds
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A;Accession: B94433
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      %1, 15591-15595, 1986
inking site in fibrinogen :
er: A92565; MUID:87057190;
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for alpha-2-plasmin inhibitor PMID:2877981
                                                                                                                                                                                                                                                                                                    PMID:6383194
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GPRVVERHQSA

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fibrinogen alpha chain

lain precursor, ext coagulation factor

extended

torm

RESULT D44234

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Fil-DyDmain: signal sequence #status predicted <SIGs Figo-629/Product: fibrinogen alpha chain #status experimental <MATS Figo-629/Product: fibrinogen alpha chain #status experimental <APTS Figo-629/Product: fibrinogen alpha chain #status experimental <APTS Figo-629/Product: fibrinogen dispha chain #status experimental <APTS Figo-629/Product: fibrin alpha chain #status experimental #status experiment
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Description: fibrinogen cleaved by thrombin yields monomers that are polymerized A;Pathway: blood coagulation C;Superfamily: fibrinogen alpha chain; fibrinogen disulfide ring homology C;Keywords: alternative splicing; blood coagulation; coiled coil; glycoprotein; liv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C; Function;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Map position: 4q28-4q28
A;Introns: 18/3; 60/3; 122/1
A;Note: the list of introns
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A;Contents: annotation, activity of cell attachment (R-G-D) motif
R;Kirschbaum, N.E.; Budzynski, A.Z.
J. Biol. Chem. 265, 13669-13676, 1990
A;Title: A unique proteolytic fragment of human fibrinogen containing the Aalpha COOH-t A;Reference number: A37117; MUID:90337977; PMID:2143188
A;Note: hementin, a protease from unamouslike
A;Note: hementin, a protease from unamouslike
A;Note: hementin, a protease from unamouslike
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A;Title: In vivo degradation of human fibrinogen A alpha: Detection of cleavage sites a A;Reference number: JC4334; MUID:96027996; pMID:7488058
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                                                                                                 Matches
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Blochem. Biophys. Res. Commun. 163, 1256-1264,
A;Title: The fibrinogen-derived peptide (RGDS)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               from the distal domain nodes.
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                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                   ,575,581,599/Cross-link: isopeptide
                                                                                       l Similarity
11; Conserv
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core and co
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F;591-593/Region: cell attachment (R-G-D) motif F;629-863/Domain: fibrinogen beta/gamma homology FBG5
F;22,460/Binding site: phosphate (Ser) (covalent) #status experimental
F;25-36/Cleavage site: Arg-Gly (thrombin) #status experimental
F;47/Disulfide bonds: interchain (to alpha-47) #status experimental
F;55/Disulfide bonds: interchain (to beta-95) #status experimental
F;64/Disulfide bonds: interchain (to beta-106) #status experimental
F;68/Disulfide bonds: interchain (to gamma-165) #status experimental
F;68/Disulfide bonds: interchain (to gamma-165) #status experimental
F;88/Disulfide bonds: interchain (to beta-106) #status experimental
F;88/Disulfide bonds: interchain (to beta-107) #status experimental
F;288,419/Binding site: carbohydrate (Asn) (covalent) #status absent
F;288,419/Binding site: carbohydrate (Asn) (covalent) #status absent
F;287,358/Cross-link: isopeptide (Gln) (interchain to Lys N6-amino of alpha) #stafing site: carbohydrate sexperimental
F;461-491/Disulfide bonds: #status experimental
F;527,558,575,581,599/Cross-link: isopeptide (Lys) (interchain to Lys N6-amino of alpha) #stafing site: carbohydrate (Asn) (covalent) #stafing site: carbohydrate 
RESULT 3
A05296
fibrinogen
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C;Comment: The alpha chain binds by 2-4 cross-links to the amino end of fibronectin.
C;Comment: The conversion of fibrinogen to fibrin is triggered by thrombin, which cleave ization sites responsible for the formation of the soft clot.
C;Comment: The soft clot is converted into the hard clot by factor XIIIa (fibrin-stabil: ger) and between alpha chains (weaker) of different monomers.
C;Comment: All fibrinogen chains are synthesized in the liver.
C;Comment: See PIR:FGHUA for the major splice form. It is not known whether this form in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Map position: 4q28-4q28
A;Introns: 18/3; 60/3; 122/1; 171/2
A;Note: the list of introns is incomplete
C;Complex: The fibrinogen molecule is a hexamer containing two sets of three nonidentic; ntained in the core. Two three-chain coiled coils emerge from this core and connect it to the core.
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,,Residues: 1-866 <7012-
,,Residues: 1-866 <7012-
,,Cross-references: GB:M58569; NID:g182406; PID:g182407
,,Note: neither the complete nucleic acid sequence nor ,,Accession: B44234
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              iochemistry 31, 11968-11972, 1992, Title: Carboxy-terminal-extended variant of the human Reference number: A44234; MUID:93090725; PMID:1457396; Accession: D44234
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Superfamily: human extended splice form fibrinogen alpha chain; is Keywords: alternative splicing; blood coagulation; glycoprotein; 1-19/Domain: signal sequence #status predicted cSIGs
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20-35/Product: fibrinopeptide A #status experimental <APT>
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Date: 10-Jun-1993 #sequence_revision 06-Sep-1996 #text_change 19-Jan-2001
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Best Local
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    alpha
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11; Conserv
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11968-11972, 1992
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A;Molecule type: protein
A;Molecule type: protein
A;Residues: 20-26,'Y',28-49;55-80;81-108 <HEN>
A;Residues: 20-26,'Y',28-49;55-80;81-108 <HEN>
R;Chung, D.W.; Rixon, M.W.; Davie, E.W.
in Proteins in Biology and Medicine, Bradshaw, R.A., ed., pp.309-328, in Proteins in Biology and Medicine, Bradshaw, R.A., ed., pp.309-328, in Proteins in Biology and Medicine, Bradshaw, R.A., ed., pp.309-328, in Proteins in Biology and Medicine, Bradshaw, R.A., ed., pp.309-328, in Proteins in Biology and Medicine, Bradshaw, R.A., ed., pp.309-328, in Proteins in Biology and Medicine, Bradshaw, R.A., ed., pp.309-328, in Proteins in Biology and Medicine, Bradshaw, R.A., ed., pp.309-328, in Proteins in Biology and Medicine, Bradshaw, R.A., ed., pp.309-328, in Proteins in Biology and Medicine, Bradshaw, R.A., ed., pp.309-328, in Proteins in Biology and Medicine, Bradshaw, R.A., ed., pp.309-328, in Proteins in Biology and Medicine, Bradshaw, R.A., ed., pp.309-328, in Proteins in Biology and Medicine, Bradshaw, R.A., ed., pp.309-328, in Proteins in Biology and Medicine, Bradshaw, R.A., ed., pp.309-328, in Proteins in Biology and Medicine, Bradshaw, R.A., ed., pp.309-328, in Proteins in Biology and Medicine, Bradshaw, R.A., ed., pp.309-328, in Proteins in Biology and Medicine, Bradshaw, R.A., ed., pp.309-328, in Proteins in Biology and Medicine, Bradshaw, R.A., ed., pp.309-328, in Proteins in Biology and Bradshaw, R.A., ed., pp.309-328, in Proteins in R.A., ed., pp.309-328, in Proteins in R.A., ed., pp.309-328, in Prote
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C;Date: 05-Jun-1987 #sequence_revision 10-Mar-1994 #text change 31-Mar-2000
C;Accession: A05294; A25715; A03117; A37505; A37506; A37507; B37505; C03117
R;Henschen, A.; Lottspeich, F.; Topfer Petersen, E.; Kehl, M.; Timpl, R.
in Protides of the Biological Fluids, Proc. 28th Colloq., ed. Peeters, H., I
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A;Molecule type: protein
A;Residues: 1,'D',3,'EGKQ',8-16 <OSB>
A;Residues: 1,'D',3,'EGKQ',8-16 <OSB>
C;Superfamily: fibrinogen alpha chain; fibrinogen disulfide ring homology
C;Keywords: blood coagulation; liver; phosphoprotein; plasma
F;1-16/Product: fibrinopeptide A #status experimental <APT>
F;3/Binding site: phosphate (Ser) (covalent) (partial) #status experiment;
                              A; Molecule type: prot
A; Residues: 1-19 < FOL
R; Timpl, R.; Fietzek,
                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Title: Amino acid sequence A;Reference number: A03117 A;Accession: A03117
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R;Sjoquist, J.; Blomback, B
Ark. Kemi 16, 425-436, 1960
                                                                                                                                                                A;Reference number: A37505
A;Accession: A37505
                                                                                                                                                                                                                                  A;Molecule type: protein
A;Residues: 1-19 <570>
R;Folk, J. E.; Gladner, J.A.; Levin,
J. Biol. Chem. 234, 2317-2320, 1959
A;Title: Thrombin-induced formation
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A; Accession: A25715
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A;Accession: A03118
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Accession: A05294
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       fibrinogen alpha chain - bovine (fragments)
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A;Residues: 1-16 <BLO>
R;Osbahr Jr., A.J.; Colma
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Best Local
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. 19, 1789-1791,
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Acta 490,
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                                  P.P.;
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81.8%;
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                                                                                                                                                                                                                                               formation
Wachter, 420-429,
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Pred. No. 0.018;
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E.;
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A;Title: Isolation of cDNA clone encoding human homologue of senescence marker protein-A;Reference number: I52491; MUID:96004897; PMID:7548213
A;Accession: S60035
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A;Title: Amino acid sequences of portions of the alpha a A;Reference number: A37507; MUID:79164394; PMID:434821
A;Accession: A37507
                                                                                                                                                                                                                                                                                                                RESULT 6
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A; Residues: 1-299 < FUJ>
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,ACCESSIUL TYPE: DNA
,MOLECULE TYPE: DNA
,RESIDUES: 1-550 <CRA>
,RESIDUES: 1-550 <CRA>
,Blombaeck, M.; (
,Blombaeck, B.; Blombaeck, M.; (
                                                                                 ;Crabtree, G.R.; Comeau, C.M.; Fowlkes, D.M.; Fornace Jr., A.J.; Malley, J.D.; Mol. Biol. 185, 1-19, 1985
. Mol. Biol. 183, 1-19, 1985
.Title: Evolution and structure of the fibrinogen genes. Random insertion of in .Reference number: A92915, MUID:86011580; PMID:4046033
                                                                                                                                                                                                                                                                      ibrinogen alpha chain precursor - rat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cross-references: GDB:9955055
Map position: Xp11.3-Xp11.2
Superfamily: senescence marker
                                                                                                                                                                                                                                                   Contains: fibrinopeptide A
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                                                                                                                                                                                      Date: 30-Jun-1987 #sequence revision Accession: A92915; B03118; A03119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cross-references: EMBL:D31815;
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                                                                                                                                                                                                                              Species: Rattus norvegicus (Norway rat)
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        Groendahl, N.J
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Pred. No. 11;
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                                                                                                                                                             submitted to the EMBL Data Library, February 1997 A;Description: The sequence of C. elegans cosmid A;Reference number: Z21526
                                                                                                                                                                                                                                                                hypothetical protein K11H12.1 - Caenorhabditis elegans C;Species: Caenorhabditis elegans C;Date: 29-Oct-1999 #sequence_revision 29-Oct-1999 #te
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                                            A; Experimental source:
                                                                                 A;Residues: 1-108 <BRA>
                                                                                                  A; Molecule type: DNA
                                                                                                                       A; Status: preliminary; translated
                                                                                                                                            A; Accession: T34436
                                                                                                                                                                                                                          R; Bradshaw, H.
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A;Moleculé type: mRNA
A;Residues: 1-336 <LES>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ş
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Title: TWIK-1, a ubiquitous human weakly A; Reference number: S65566; MUID: 96183184;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EMBO J. 15, 1004-1011, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  R;Lesage, F.; Guillemare, E.; Fink, M.; Duprat, F.; Lazdunski, M.; Romey, G.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  inward rectifier potassium channel TWIK-1 - human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                F;404-434/Disulfide bonds:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              F;20-36/Product: fibrinopeptide A #status predicted <510>
F;20-36/Product: fibrinopeptide A #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Residues: 20-29 <BLO>
C;Comment: Unlike the beta and
C;Comment: Fibrinogen chains ar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Title: Studies on fibrinopeptides from mammals A; Reference number: A03118
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C;Superfamily:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Introns: 18/3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C; Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Molecule type:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Accession:
                                                                                                                                                                                                                                               Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ;37-550/Product: fibrinogen alpha chain #status predicted;58-186/Domain: fibrinogen disulfide ring homology <FDR>;48/Disulfide bonds: interchain (to alpha-48) #status pred
                                                                 Cross-references:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ;Species: Homo sapiens (man)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Superfamily: fibrinogen alpha chain; fibrinogen disulfide; Keywords: blood coagulation; liver; plasma; platelet; 1-19/Domain: signal sequence #status predicted <SIG>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 .81/Disulfide bonds: interchain (to gamma-165)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 3 RVVERHQSA 11
                                                                                                                                                                                                                                                 T34436
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            B03118
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                 EMBL: U88168;
                                                                                                                                                                                                                                                              #sequence_revision
                                                                                                                                                                                                                                                                                                                                                                                                           20
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77.8%;
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are synthesized
                                       58; PIDN:AAC24398.1; GSPDB:GN00022; CESP:K11H12.1 Bristol N2; clone K11H12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      171/3
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Pred. No. 21;
1; Mismatches
                                                                                                                         from
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                                                                                                                         GB/EMBL/DDBJ
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21;
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Gaps

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channel with a novel

Barhanin

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C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 02-Mar-2001 #sequence_revision 02-Mar-2001 #text_change 31-Mar-2001
C;Accession: F96769
R;Theologis, A.; Ecker, J.R.; Palm, C.J.; Federspiel, N.A.; Kaul, S.; White, O Chin, C.W.; Chung, M.K.; Conn, L.; Conway, A.B.; Conway, A.R.; Creasy, T.H.; ansen, N.F.; Hughes, B.; Huizar, L.
Nature 408, 815-820, 2000
A;Authors: Hunter, J.L.; Jenkins, J.; Johnson-Hopson, C.; Khan, S.; Khaykin, E.C.A.; Li, J.H.; Li, Y.; Lin, X.; Liu, S.X.; Liu, S.A.; Luros, J.S.; Maiti, R.; Rizzo, M.; Rooney, T.; Rowley, D.; Sakano, H.
A;Authors: Salzberg, S.L.; Schwartz, J.R.; Shinn, P.; Southwick, A.M.; Sun, H.
ker, M.; Wu, D.; Yu, G.; Fraser, C.M.; Venter, J.C.; Davis, R.W.
A;Title: Sequence and analysis of chromosome 1 of the plant Arabidopsis.
A;Accession: F96769
                                                                                                    Science 277, 1453-1462, 1997
A; Title: The complete genome sequence of Escherichia coli K-12.
A;Reference number: A64720; MUID:97426617; PMID:9278503
A;Scression: H65094
A;Starum: No. 272-278504
                                                                                                                                                                                                                                                                                                                    C;Date: 20-Sep-1984 #sequence_revision 20-Sep-1984 #text_change C;Accession: A03423; H65094; A03422; I77532
R;Burton, Z.F.; Gross, C.A.; Watanabe, K.K.; Burgess, R.R.
Cell 32, 335-349, 1983
A;Title: The operon that encodes the sigma subunit of RNA polyme A;Reference number: A02749; MUID:83129424; PMID:6186393
A;Accession: A03423
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Map position: 4
A; Introns: 22/3; 61/
C; Superfamily: bolA
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A;Residues: 1-581 <BLAT>
A;Cross-references: GB:AE000388; GB:U00096; NID:g1789441;
A;Experimental source: strain K-12, substrain MG1655
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Map position: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Cross-references:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Residues: 1-552 <STO>
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A; Residues: 1-581 <BUR>
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                                                                      A; Molecule type: DNA
                                                                                                                                                                                                      Cross-references: GB:J01687; NID:g147753; PIDN:AAA24600.1; PID:g147755; Blattner, F.R.; Plunkett III, G.; Bloch, C.A.; Perna, N.T.; Burland, V.; Riley, A.; Rose, D.J.; Mau, B.; Shao, Y.
                                                                                           Status: preliminary; nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Species: Escherichia coli
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GPRVREAHSAA 131
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                                                                                               acid
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Pred. No.
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                                                                                           sequence
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                            PIDN:AAC76102.1; PID:g1789447
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.S.; Maiti, R.;
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DNA Res. 8, 11-22, 2001
A;Title: Complete genome sequence of enterohemorrhagic Escherichia coli O157:H7
A;Reference number: A99629; MUID:21156231; PMID:11258796
                                                                   gasawara, N.; Yasunaga,
                                                                                        C; Accession: E91122
R; Hayashi, T.; Makir
                                                                                                                              C;Species: Escherichia coli
C;Date: 18-Jul-2001 #sequence_revision 18-Jul-2001 #text_change 27-Nov-2001
                                                                                                                                                                                                                                                                                               밁
                                                                                                                                                                                   DNA primase [imported] - Escherichia coli (strain O157:H7, substrain RIMD 0509952)
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Makino, K.; Ohnishi,

T.; Kuhara,

M.; Kurokawa, F S.; Shiba, T.;

K.; Ishii, Hattori,

Shinagawa,

K.; Han, H.

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and

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A; Map position: 67 min C; Superfamily: DNA primase C; Keywords: DNA replication
                               S
                                                                                                                                                                                                                                                                                      A;Title: Genome sequence of enterohemorrhagic Escherichia coli A;Reference number: A85480; MUID:21074935; PMID:11206551 A;Accession: D85967
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       R;Smiley, B.L.; Lupski, J.R.; Svec, P.S.; McMacken, R.; Proc. Natl. Acad. Sci. U.S.A. 79, 4550-4554, 1982 A;Title: Sequences of the Escherichia coli dnaG primase A;Reference number: A03422; MUID:83014926; PMID:6750604
                                                                                                                                             C;Superfamily:
                                                                                                                                                                                                                A;Molecule type: DNA
A;Residues: 1-581 <STO>
A;Cross-references: GB:
                                                                                                                                                                                                                                                                                                                                              iller, L.; Grotbeck, E.J.
Nature 409, 529-533, 2001
                                                                                                                                                                                                                                                                                                                                                             R;Perna, N.T.; Plunkett III, G.; Burland, V.; Mauiller, L.; Grotbeck, E.J.; Davis, N.W.; Lim, A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Gene: dnaG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Cross-references: EMBL:V00346; NID:g42867; PIDN:CAA23636.1; C;Comment: This enzyme, an RNA polymerase, interacts with DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Molecule type: DNA
A;Residues: 1-23,'N',25-81 <RES>
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                                                                       Matches
                                                                                      Query Match
Best Local
                                                                                                                                                                                                 Experimental source:
                                                                                                                                                                                                                                                                         Status: preliminary;
                                                                                                                                                                                                                                                                                                                                                                                                   ;Date: 16-Feb-2001 #sequence_revision 16-Feb-2001 #text_change 27-Nov-2001
;Accession: D85967
                                                                                                                                                                                                                                                                                                                                                                                                                                         ;Species: Escherichia
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Best Local
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ol. Gen. Genet. 189, 48-57, 1983
104
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1-23,'N',25-105,'XPDRAPSEANALSVDGRSEYVLPTIFTTTCCHVCAPVSG',145-156,'GLVLR',1
                                                                                        Similarity
GPSQIERHQ 112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GPSQIERHQ 112
                                 GPRVVERHQ 9
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                                                                                                                                           DNA primase
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                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA primase [imported] - Escherichia coli (strain O157:H7,
ichia coli
                                                                                                                                                                                               GB:AE005174; NID:g12517648; PIDN:AAG58200.1; GSPDB:GN00145; UWGP:Z4
ce: strain O157:H7, substrain EDL933
                                                                                      66.7%;
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                                                                                      Pred. No. 55;
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Pred. No.
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A.; Dimalanta,
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lanta, E.;
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K.; Apodaca
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A;Description: calcium binding A;Note: age-associated decrease C;Superfamily: senescence marker C;Keywords: calcium binding
                                                                                                                                                                                                                                                                                                         R; Fujita, T.; Shirasawa, T.; Maruyama, N. Biochim. Biophys. Acta 1308, 49-57, 1996 A; Title: Isolation and characterization of
                                                                                                                                                                                                                   A;Molecule type: mRNA
A;Residues: 1-299 <FUJ>
A;Cross-references: EMBL:U28937; NID:gl143999; PIDN:AAC52721.1;
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\;Residues: 1-54 <FUW>
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A;Variety: PCC 6716
C;Date: 22-Jan-1994 #sequence_revision 12-Apr-1996 #text_change 08-Oct-1999
C;Accession: S36971; S31895
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S36971
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A;Title: Organization and sequences of genes for the subunits of
A;Reference number: 836960; MUID:93371369; PMID:8363578
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          R;van Walraven, H.S.; Lut
Biochem. J. 294, 239-251,
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C;Superfamily:
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A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-581 <HAY>
                                                       Superfamily: senescence marker;Keywords: calcium binding
                                                                                                                                  ;Cross-references: EMBL:U32170; NID:g1144337; PIDN:AAD03478.1;
;Experimental source: liver
                                                                                                                                                                                                                                                                                                                                                         ;Species: Mus musculus (house mouse);Date: 04-Dec-1997 #text_change 20-Sep-1999;Accession: S72173; S72174
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A;Experimental source: strain O157:H7, substrain RIMD 0509952
C;Genetics:
                                                                                                                                                                                                                                                                                 Reference number: $72173; MUID:96328264; PMID:8765750
            Best Local Similarity
                                Query Match
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61.4%;
60.0%;
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                                                                           protein-30
        Score 35;
Pred. No.
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Pred. No. 9.3;
4; Mismatches
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Pred. No.
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        DB 2;
44;
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                   Length 299;
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Search completed: June 27, 2003, 17:18:35 Job time: 16.3 secs

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                                                                                                                        A;Status: preliminary
A;Molecule type: mRNA
A;Molecule type: mRNA
A;Residues: 1-147,'D',149-299 <FUU>
A;Cross-references: EMBL:X69021; NID:g57254; PIDN:CAA48786.1; PID:g57255
C;Superfamily: senescence marker protein-30
C;Keywords: calcium binding
                                                                                                                                                                                                                                          A;Residues: 84-299 < SH2>
A;Cross-references: EMBL:D14327
A;Cross-references: EMBL:D14327
R;Fujita, T; Shirasawa, T.; Uchida, K.; Maruyama, N.
Biochim. Biophys. Acca 1132, 297-305, 1992
A;Title: Isolation of cDNA clone encoding rat senescence marker protein-30 (SMP30) and A;Reference number: S27203; MUID:93041931; PMID:1420310
A;Accession: S27203
                                                                                                                                                                                                                                                                                                                                                               A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Title: Molecular cloning and sequencing of A;Reference number: S34588; MUID:93351639; A;Accession: S34588
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C;Accession: S34588; S34589; R;Shimokawa, N.; Yamaguchi, M.
FEBS Lett. 327, 251-255, 1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C;Species: Rattus norvegicus (Norway rat)
C;Date: 02-Dec-1993 #sequence revision 10
C;Accession: S34588; S34589; S27203
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1;Residues: 1-299 <SHI>
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     124
                                2 PRVVERHOSA 11
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                                                                                      Similarity
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PAVLERHQGS 133
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                                                                                  61.4%;
                                             Pred. No.
2: Mismatches
                                                                              Score 35;
Pred. No.
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44;
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     Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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57
1 GPRVVERHQSA 11
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Match
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P02673; P14464;

21-JUL-1986 (Rel. 0

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                                                                                                                                                              Blood coagulation; P
PEPTIDE 1
MOD_RES 3
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NON_TER 28
SEQUENCE 28 AA; 7
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Osbahr A.J. Jr., Colman R.W., Laki K., Gladner J.A.;
"The nature of the peptides released from canine fit Biochem. Biophys. Res. Commun. 14:555-558(1964).
-I- FUNCTION: FIBRINGEN HAS A DOUBLE FUNCTION: YIEL POLYMERIZE INTO FIBRIN AND ACTING AS A COFACTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SPECIES=C.familiaris, and V.vulpes;
Blomback B., Blomback M., Grondahl N.J.;
"Studies on fibrinopeptides from mammals.";
Acta Chem. Scand. 19:1789-1791(1965).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE=76081726; PubMed=1198547;
Birken S., Wilner G.D., Canfield R.E.;
"Studdes of the structure of canine fibrinogen.";
Thromb. Res. 7:599-610(1975).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalla; Eutheria; Carnivora; Fissipedia; Canidae; NCBI_TaxID=9615, 9627;
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Vulces vulpes (Red
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A05296;
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41, Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Folk J.E., Gladner J.A., Levin Y., Folk W.F., Gladner J.A., Levin Y., Thrombin-induced formation of co-fibrin. III. Acid studies and summary of sequential evidence on peptic J. Biol. Chem. 234:2317-2320(1959).
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Sjoquist J., Blomback B., Walle Sjoquist J., Blomback b., Walle "Amino acid sequence of bovine Ark. Kemi 16:425-436(1960).
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                                                                                                   MEDLINE=20087206; PubMed=10618375;
Brown J.H., Volkmann N., Jun G., Henschen-Edman A.H., C.
"The crystal structure of modified bovine fibrinogen.";
Proc. Natl. Acad. Sci. U.S. A. 97.85-90(2000).
-i- FUNCTION: FIBRINOGEN HAS A DOUBLE FUNCTION: YIELDIN POLYMERIZE INTO FIBRIN AND ACTING AS A COFACTOR IN
                                                                                                                                                                                                                                              SEQUENCE OF 20-49; 243-268 AND 553-580.
Henschen A., Lottspeich F., Topfer-Petersen E.,
                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=77112616; PubMed=836881;
Timpl R., Fietzek P.P., Wachter B.,
"Disulfide-linked cyanogen bromide
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Murakawa M.;
                                                                                                                                                                                 (-RAY CRYSTALLOGRAPHY (4.0 ANGSTROMS)
SUBUNIT: HEXAMER CONTAINING 2 SETS OF 3 NONIDENTICAL CHAINS (ALPHA, BETA AND GAMMA), LINKED TO EACH OTHER BY DISULFIDE BOND MISCELLANGOUS: CONVERSION OF FIBRINGGEN TO FIBRIN IS TRIGGERED THROMBIN, WHICH CLEAVES FIBRINOPEPTIDES A AND B FROM ALPHA & BE CHAINS, AND THUS EXPOSES THE N-TERMINAL POLYMERIZATION SITES RESPONSIBLE FOR THE FORMATION OF THE SOFT CLOT. THE SOFT CLOT CONVERTED INTO THE HARD CLOT BY FACTOR XIIIA WHICH CATALYZES TH
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J D.W., Rixon M.W., Davie E.

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Oxford (1980).
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an email to license@isb-sib.ch).
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                                                           36, Created)
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41, Last annotation updat
protein-30 (SMP-30) (Regu
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158
194
432
523
                                  Chordata;
Primates;
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170
208
220
353
439
                                                                                                                                                                                                                                                                                                                                                                                     Plasma;
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                                                                                                                                                                       80.7%;
81.8%;
                                                                                                                                                                                                                                                                                         52
                                                                                                                                                                                                                                                                                                                                                                       Glycoprotein; Coiled coil; 3D-structure, FIBRINOPEPTIDE A.
FIBRINOGEN ALPHA CHAIN.
                                                                                                                                                                                                       BY SIMILARITY.

R -> K (IN REF. 1).

IL -> LI (IN REF. 1).

RE -> KK (IN REF. 1).

MISSING (IN REF. 1).

KV -> S (IN REF. 2).
                                                                                                                                                                                                                                                                                                        INTERCHAIN (WITH C-4 INTERCHAIN (WITH C-5 SIMILARITY).
                                                                                                                                                                       Score 46;
Pred. No.
                                    Craniata; Vertebrata; Euteleostomi;
                               Catarrhini;
                                                                                                                                                                                                                                                                     SIMILARITY).
INTERCHAIN (WITH C-165 IN GAMMA)
                                                                                                                                                                                                                                                                  SIMILARITY).
                                                                                                                                                                                                                                                                                     SIMILARITY).
INTERCHAIN (WITH
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COILED COIL (POTENTIAL)
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                                                           (Regucalcin) (RC)
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                                Hominidae;
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Usage
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This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         - HUMAN
-Y711 HUMAN
094819;
16-OCT-2001
16-OCT-2001
                                                                                                                               Nagase T., Ishikawa K.-I., Suyama M., Kikuno R., Miyajima Tanaka A., Kotani H., Nomura N., Ohara O., "Prediction of the coding sequences of unidentified human The complete sequences of 100 new cDNA clones from brain v for large proteins in vitro.", DNA Res. 5:277-286(1998)
                                                                                 -i- SIMILARITY: CONTAINS 1 BTB/POZ DOMAIN.
                                                                                                                                                                                                                                                      TISSUE=Brain;
MEDLINE=99087487; PubMed=9872452;
                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                  Hypothetical KIAA0711.
                                                                                                                                                                                                                                                                                                                                        NCBI_TaxID=9606,
                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Genew; HGNC: 9
MIM; 300212;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Calcium-binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EMBL; AB028125; BAA78693.1; -. EMBL; AB032064; BAA84082.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 use by non-profit institutions as long modified and this statement is not removed. entities requires a license agreement (See or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          .6-OCT-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       regucalcin.",
Int. J. Mol. Med. 5:283-287(2000).
Int. TUNCTION: MAY PLAY A ROLE IN THE REC.
IN THE LIVER. DECREASE OF SMP30 LEAL
CALCIUM SIGNALING IN THE AGED LIVER.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=96004897; PubMed=7548213; Fujita T., Mandel J.L., Shirasawa T., Hino O., Shirai T., Maruyama "Isolation of cDNA clone encoding human homologue of senescence marker protein-30 (SMp30) and its location on the X chromosome.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   between
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "Transcript heterogeneity of the human gene for Ca2+-binding protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Misawa H., Yamaguchi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PubMed=10677570;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   s SWISS-PROT entry is copyright. It is produced through a collaboration ween the Swiss Institute of Bioinformatics and the EMBL outstation European Bioinformatics Institute. There are no restrictions on it by non-profit institutions as long as its content is in no wa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SUBCELLULAR LOCATION: Cytoplasmic (ESIMILARITY: BELONGS TO THE SMP-30 /
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PAVLERHQGA 133
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(Rel. 40, Last sequence update)
(Rel. 40, Last annotation updat)
protein KIAA0711
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 38;
Pred. No. 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  95BA1C73B7B77635 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               update)
                                                                                                                                                              unidentified human genes. XI clones from brain which code
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                noved. Usage by and for commercial (See http://www.isb-sib.ch/announce)
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SEQUENCE OF 20-36.

Blomback M., Grondahi N. ...,

"Studies on fibrinopeptides from mammals.";

"Studies on fibrinopeptides from mammals.";

"Studies on fibrinopeptides from mammals.";
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01-OCT-1996
16-OCT-2001
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                                                                                                                 MEDLINE=87134033; PubMed=3817019; Sobczak J., Lotti A.-M., Taroux P., Du "Molecular cloning of mRNA sequences t liver regeneration."; Exp. Cell Res 169:47-56(1987). Exp. Cell Res 169:47-56(1987). PURCTION: FIBRINGEN HAS A DOUBLE POLYMERIZE INTO FIBRIN AND ACTING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus MCBI_TaxID=10116;
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P06399;
                                                                                                                                                                                                                                                                                                                                                       SEQUENCE OF 458-550 FROM N.A. (ALPHA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Malley J.D.,
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Submitted (
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01-OCT-1996 (Rel. 34, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Fibrinogen alpha/alpha-E chain precursor (Contains: Fibrinopeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE
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InterPro; IPR001798;
Pfam; PF00651; BTB;
                                                                                                                                                                                                                                                                                                                                 STRAIN=Wistar;
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   AGGREGATION.
SUBUNIT: HEXAMER CONTAINING 2
(ALPHA, BETA AND GAMMA), LINKE
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(MAR-1995) to t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A
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72.7%;
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KELCH 2.

KELCH 3.

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KELCH 4.

KELCH 4.
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EMBL/GenBank/DDBJ databases
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
      ARE
                                 ED TO
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TS OF 3 NONIDENTICAL TO EACH OTHER BY DISC
                                                                                                                                                                                                                                         transiently induced
                                                                                                                                                                                                                                                                                                                                                          FORM)
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                                                             CHAINS
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CONFLICT
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SMART; SM00186; FBG; 1.
PROSITE; PROAFT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                entities
or send a
                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL; X86561; CAA60264.1;
EMBL; X86561; CAA60263.1;
EMBL; M35601; AAA41158.1;
PIR; A03119; FGRTA.
                                                                                                                                                                                                                                                                                                                                                               Blood
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              <u>-</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              is SWISS-PROT entry is copyright. It is produced through a collaboration tween the Swiss Institute of Bioinformatics and the EMBL outstation - se European Bioinformatics Institute. There are no restrictions on its e by non-profit institutions as long as its content is in no way dified and this statement is not removed. Usage by and for commercial titles requires a license agreement (See http://www.isb-sib.ch/announce/send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PIW: THE ALPHA CHAIN IS NOT GLYCOSYLATED.

MISCELLANEOUS: CONVERSION OF FIBRINGGEN TO FIBRIN IS TRIGGERED BY THROMBIN, WHICH CLEAVES FIBRINOFEPTIDES A AND B FROM ALPHA & BETA CHAINS, AND THUS EXPOSES THE N-TERMINAL POLYMERIZATION SITES RESPONSIBLE FOR THE FORMATION OF THE SOFT CLOT IS CONVERTED INTO THE HARD CLOT BY FACTOR XIIIA WHICH CATALYZES THE EPSILON-(GAMMA-GLUTAMYLLYSINE CROSS-LINKING BETWEEN GAMMA CHAINS (STRONGER) AND BETWEEN ALPHA CHAINS (WEAKER) OF DIFFERENT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DIVERGING FROM THIS NODULE ARE 2 THREE-CHAIN COILED COILS, WHICH CONNECT THE CENTRAL NODULE TO THE DISTAL NODULES CONTAINING THE DISTAL DOMAINS. EXTENDING FAR PERIPHERALLY ARE THE LONG CARBOXYL ENDS OF THE ALPHA CHAINS.

ALTERNATIVE PRODUCTS: 2 ISOFORMS; ALPHA-E AND ALPHA; ARE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SIMILARITY: CONTAINS 1 FIBRINOGEN C-TERMINAL DOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PREDOMINANT FORM
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37
                                             . Similarity 7; Conserv
                     GPRVVERHOS
GPRIVERQPS
                                             Conservative
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404
547
551
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212
270
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181
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55
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRODUCTS: 2 ISOFORMS; ALPHA-E AN ALTERNATIVE SPLICING. THE ALPHA
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434
550
782
782
140
212
276
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181
46
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                                                         66.7%;
70.0%;
                                                                                                                                                                                                                                                                                                                                                                lasma;
                                                                                              MW;
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Q -> E (IN REF.
D -> E (IN REF.
                                                         Score 38; DB
Pred. No. 11;
                                                                                                        ASRGDLP -> LREEIYO (IN REF. S -> K (IN REF. 4).
                                                                                                                                                                   DCDD -> GIHA (IN ISOFORM ALPHA)
MISSING (IN ISOFORM ALPHA)
                                                                                                                                                                                                                                                              INTERCHAIN (WITH C-48') (BY SIMILARITY).
INTERCHAIN (WITH BETA) (BY SIMILARITY).
INTERCHAIN (WITH C-49 IN GAMMA)
                                                                                                                                                                                                                                                                                                               FIBRINOGEN
CLEAVAGE (I
                                                                                                                                                                                                                                                                                                     FIBRINOPEPTIDE
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                                                                                                                                                                                                                    (BY SIMILARITY)
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                                                                                            744834DAE76D34C2
                                              Mismatches
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                                                                                                                                           DEGAG (IN REF. 3).
IN REF. 2).
                                                                                                                                                                                                                                                                                                              N ALPHA-E CHAIN. (BY THROMBIN; RE
                                                                     1;
                                                                                                                                                                                                                                                                                                                                                                Alternative
                                                                                                                                                                                                                            ( BETA)
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                                                                                                                                                                                                       BETA) (BY SIMILARITY)
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                                                                     Length 782
                                                                                              CRC64;
                                              Indels
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                                                                                                                                                                                                                                                                                                                 RELEASE
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GAMMA)
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RESULT 7
CIWI HUMAN
ID CIWI HUMAN STANDARD;
AC 000180; Q13307;
AC 16-OCT-2001 (Rel. 40, Created)

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Orias M., Velazquez H., Tung F., Lee G., Desir G.V.;

"Cloning and localization of a double-pore K channel, KCNK1: exclus expression in distal nephron segments.";

Am. J. Physiol. 273:F663-F666 (1997).

-!- FUNCTION: WEAKLY INWARD RECTIFYING POTASSIUM CHANNEL.

-!- SUBURIT: HOMODIMER (POTENTIAL).

-!- SUBCRILULAR LOCATION: Integral membrane protein (Potential).

-!- SUBCRILULAR LOCATION: Integral membrane protein KPRELS IN HEART EXPRESSED WITH HIGH LEVELS IN HEART BRAIN AND LOWER LEVELS IN PLACENTA, LUNG, LIVER AND KINNEY.

-!- MISCELLANEOUS: INHIBITED BY BARIUM, QUININE, QUININE AND INTERNAL ACIDIFICATION. ACTIVATED BY PROTEIN KINASE C.

-!- SIMILERITY: BELONGS TO THE TWO PORE DOMAIN FAMILY OF POTASSIUM.
                                                                                                                                                                                                                                                             InterPro; IPR003280; K+channel 2pore.
InterPro; IPR001622; K+channel pore.
InterPro; IPR000636; M+channel nlg.
InterPro; IPR001779; TWIK1 channel.
                                                                                                                                                                                                                                                                                                                                                                MIM;
                                                                                                                                                                                                                                                                                                                                                                                                   EMBL; U76996; AAB97878.1; -. EMBL; U90065; AAB51147.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                         EMBL; U33632; AAB01688.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EN the European Bioinformatics Institute. There are no resti
                                            TRANSMEM
                                                                                     TRANSMEM
                                                                                                                                                                     Glycoprotein
                                                                                                                                                                                                                                                      Pfam; PF00520;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     entities requires a license agreement (S or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Goldsrein S.A.N., Wang K.-W., Ilan N., Pau "Sequence and function of the two P domain implications of an emerging superfamily.", J. Mol. Med. 76:13-20(1998).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EMBO
  TRANSMEM
                                                             DOMAIN
                                                                                                           DOMAIN
                                                                                                                               TRANSMEM
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                                                                                                                                                                                                                                                                                                                                                                                    Genew;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            use by non-profit institutions as lor modified and this statement is not remove
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=98026667; PubMed=9362344;
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                                                                                                                                                                                           Ionic channel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=98122696;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TISSUE=Brain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         novel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "TWIK-1, a ubiquitous human weakly inward rectifying K+ channel with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       channel protein TWIK-1) (PCKCNK1 OR TWIK1 OR HOHO1 OR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16-OCT-2001 (Rel. 40, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Potassium channel subfamily K member 1 (Inward rectifying potassium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Barhanin J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE=96183184; PubMed=8605869;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  lesage F.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              l structure.";
J. 15:1004-1011(1996)
                                                                                                                                                                                                                                                                                                                                                                               HGNC:6272; KCNK1.
                                                                                                                                                                                                               PR01096;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Guillemare
                                                           21
104
133
154
                                                                                                                                                                                                                              ion_trans; 1.
3; 2POREKCHANEL.
                                                                                                                                                                                      Transmembrane;
                                                                                                                                                                                                            TWIKICHANNEL.
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41
130
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POTENTIAL
               PORE-FORMING
                                     POTENTIAL
                                                         CYTOPLASMIC (POTENTIAL)
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P domain potassium channels:
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MBL outstation -
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RESULT 8
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Best Local S
Matches 7
                                                                                                                                                                                                                                                                                                        mouse twik K+ channel gene.";
FEBS Lett. 425:310-316(1998).
-!- FUNCTION: WEAK INWARDLY RECTIFYING POTASSIUM CHANNEL.
-!- SUBUNIT: HOMODIMER (POTENTIAL).
                                                                                                                                                                                                                                                                                                                                        STRAIN=129/SvJ; TISSUE=Liver;
MEDLINE=98218573; PubMed=9559671;
Arrighi I., Lesage F., Scimeca J.-C., Carle G.F., Barhanin J.
"Structure, chromosome localization, and tissue distribution mouse twik K+ channel gene.";
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CARBOHYD
MUTAGEN
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A
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Lesage F., Lauritzen I., Duprat F.,
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Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          channel protein
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16-OCT-2001 (Rel. 40, Last sequence update).
16-OCT-2001 (Rel. 40, Last annotation update)
Potassium channel subfamily K member 1 (Inward
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=10090;
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                                                                                                                                                                                                         SUBCELLULAR LOCATION: Integral membrane protein (Potential).
TISSUE SPECIFICITY: WIDELY EXPRESSED HIGHEST EXPRESION IN BRA
KILNEY, TYRROID, SALIVARY GLAND, ADRENAL GLAND, PROSTATE,
EPIDIDYMIS, UTERUS, PLACENTA, COLON AND JEJUNUM. MODERATE
EXPRESSION IN EYES, PITUITARY, PANCREAS, SMOOTH MUSCLE, TESTIS
OVARY, VERY LOW LEVELS IN LUNG, AORTA, LIVER, HEART, SKELETAL
MUSCLE, THYMUS AND SPLEEN. IN THE BRAIN, HIGHEST EXPRESSION IN
CEREBELLAR GRANULE CELLS, BRAINSTEM, HIPPOCAMPUS AND CEREBRAL
                                                                                                                                                               CONCEPTION. EXPRESSION STABILIZES AFTER DAY 8
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77.88;
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T->A: NO EFFECT ON CHANNEL ACTIVITY
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EMBL; AF033017;

AAC16973.1;

-i- MISCELLANEOUS: INHIBITED BY QUININE, ACIDIFICATION. ACTIVATED BY PROTEIN -i- SIMILARITY: BELONGS TO THE TWO PORE

KINASE C.
DOMAIN FAMILY OF , BARIUM, KINASE C

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15-JUL-1998 (Rel. 3
16-OCT-2001 (Rel. 4
Hypothetical 12.1)
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P91375;
15-JUL-1998
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                                                                                                                                                           This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EN the European Bioinformatics Institute. There are no restruse by non-profit institutions as long as its content modified and this statement is not removed. Usage by and
                                                                                                                                                                                                                                                                                        Caenorhabditis elegans.
Eukaryota; Metazoa; Nem
Rhabditidae; Peloderina
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SEQUENCE
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InterPro; IPR001622; K+channel_pore.
InterPro; IPR001636; M+channel_nlg.
InterPro; IPR001779; TWIKI channel.
Pfam; PF00520; ion_trans; I.
                                                                                                                                                                                                                     Submitted (FEB-1997) to -!- SIMILARITY: BELONGS
                                                                                                                                                                                                                                         Bradshaw H.
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                                                                                                              EMBL; U88168; AAC24398.1; -. WormPep; K11H12.1; CE12146.
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STRAIN-Bristol N2;
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Pfam; PF01722; BolA; 1.
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PRO1096; TWIK1CHANNEL.
                                                                                                                                          s requires a license agreement (See an email to license@isb-sib.ch).
                                    Similarity 6; Conser
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protein K11H12.1 in chr
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CYTOPLASMIC (POTENTIAL)
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MEDLINE=83014926; PubMed=6750604; Smiley B.L., Lupski J.R., Svec P.S. "Sequences of the Escherichia coli of its expression."; Proc. Natl. Acad. Sci. U.S.A. 79:45
                                                                                                                                                                                                                                                                                              Iida T., Ta
Kuhara S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rose D.J., Mayhew G.F., Evans P.S., Gregor J., Kirkpatrick H.A., Posfai G., Hackett J., Klink S., Boutin A., Shao Y., Miller L., Grotbeck E.J., Davis N.W., Lim A., Dimalanta E.T., Potamousis K., Apodaca J., Anantharaman T.S., Lin J., Yen G., Schwartz D.C., Welch R.A., Blattner F.R.;
"Genome sequence of enterohaemorrhagic Escherichia coli O157:H7."
Nature 409:529-533(2001).
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21-JUL-1986
21-JUL-1986
15-JUN-2002
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Biochim.
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                                                                                                                                                                                                                                                                                                                                     STRALN=0157:H7 / RIMD 0509952;
MEDLINE=21156231; PubMed=11258796;
Hayashi T., Makino K., Ohnishi M., F
Han C.-G., Ohtsubo E., Nakayama K.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Blattner F.R., Plunkett G. III, Bloch C.. Riley M., Collado-Vides J., Glasner J.D. Gregor J., Davis N.W., Kirkpatrick H.A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=21074935; PubMed=11206551;
Perna N.T., Plunkett G. III, Burland V.,
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MEDLINE=97426617;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=83129424; PubMed=6186393;
MEDLINE=83129424; PubMed=8186393;
MEDLINE=8312944; PubMed=8186393;
MEDLINE=8312944; PubMed=8186393;
MEDLINE=8312944; PubMed=818644;
MEDLINE=831294; PubMed=818644;
MEDLINE=831294; PubMed=818644;
MEDLINE=831294; PubMed=81864; PubMed=81864
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Stamford N.P.J., 1
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STRAIN=0157:H7 / I
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                   (-RAY CRYSTALLOGRAPHY (2.9 ANGSTROMS)
                                                                                                                                                                                                                                                ashi T., Makino K., Ohnishi M., Kurokawa K., Ishii K., Yokoy C.-G., Ohtsubo E., Nakayama K., Murata T., Tanaka M., Tobe a T., Takami H., Honda T., Sasakawa C., Ogasawara N., Yasuna ara S., Shiba T., Hattori M., Shinagawa H.; Shiba T., Hattori M., Shinagawa H.; Title genome sequence of enterohemorrhagic Escherichia coli 7:H7 and genomic comparison with a laboratory strain K-12.";
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UL-1986 (Rel. 01, Last sequence up
UN-2002 (Rel. 41, Last annotation
primase (EC 2.7.7.-).
OR DNAP OR PARB OR B3066 OR Z441:
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Q64374; Q60944;
Q1-NOV-1997 (Rel.
Q1-NOV-1997 (Rel.
Q1-NOV-1998 (Rel.
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EMBL; V00274; CAA33531.1; ALT_FRAME.

EMBL; U28379; AAA89146.1; -...

EMBL; AE000388; AAC76102.1; -...

EMBL; AE0005536; AAG58200.1; -...

EMBL; AP002564; BAB37372.1; -...

EMBL; A03423; RYEC2.
Fujita T., Shirasawa T., Maruyama N., "Isolation and characterization of gemouse senescence marker protein-30 (S)
                                                                                                    muscutus (mouse).

Bumaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleosto
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
                                           MEDLINE=96328264; PubMed=8765750;
                                                                      SEQUENCE FROM N.A.
                                                                                                                                    RGN OR SMP30.
Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=10090;
                                                                                                                                                                  Senescence marker
                                                                                                                                                                                                                                                  MOUSE
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Pfam; PF01807; zf-CHC2; 1.
ProDom; PD002988; Znf_CHC2; 1.
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ECO2DBASE; G060.0; 6TH EDITION.
ECOCene; EG10239; dnaG.
InterPro; IPR002936; DNAprim toprim.
InterPro; IPR002694; Znf_CHC2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              J. MOI. Biol. 300:353-362(2000).

-!- FUNCTION: DNA PRIMASE IS THE POLYMERASE THAT SYNTHESIZES RNA PRIMERS FOR THE OKAZAKI FRAGMENTS ON BOTH TEMPLATE S' REPLICATION FORKS DURING CHROMOSOMAL DNA SYNTHESIS.

-!- COPACTOR: BINDS ONE ZINC ION PER MOLECULE.

-!- SUBBUNIT: MONOMER.
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Escherichia coli
topoisomerases."
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                                                     TISSUE=Liver;
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35, Last sequence update)
37, Last annotation updat
protein-30 (SMP-30) (Regu
                                                                                                                                                                                                                                                                                                                                9
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66.7%;
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Ystal structure of the catalytic
onfirms a structural link to DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DIFFERS
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Pred. No.
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 genomic an (SMP30).";
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                                                                                                                                                              (Regucalcin)
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          and cDNA clones encoding
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Best Local :
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15-JUN-2002 (Rel. 4
15-JUN-2002 (Rel. 4
15-JUN-2002 (Rel. 4
                                       Vertebrate species.";

Int. J. Mol. Med. 6:191-196(2000).

Int. J. Mol. Med. 6:191-196(2000).

Int. J. Mol. Med. 6:191-196(2000).

IN THE LIVER. DECREASE OF SMP30 LEADS TO THE DY CALCIUM SIGNALING IN THE AGED LIVER (BY SIMILAR I-- SUBCELLULAR LOCATION: Cytoplasmic (By similarit SIMILARITY: BELONGS TO THE SMP-30 / CGR1 FAMILY
     This
                                                                                                                                                                                                                  Misawa H., Yamaguchi M.; "The gene of Ca2+~binding
                                                                                                                                                                                                                                                                    MEDLINE=20351777; PubMed=10891565
                                                                                                                                                                                                                                                                                                 TISSUE=Liver;
                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RABIT
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                                                                                                                                                                                                                                                                                                                                                                                               Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Oryctolagus
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EMBL; U32170; AAD03478.1; --
EMBL; D86217; BAA13046.1; --
SWISS-2DPAGE; Q64374; MOUSE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This SWISS-PROT entry is copyright. It is produced through a continuous transfer and this statement is not removed. Usage by and for entities requires a license agreement (See http://www.isb-sib.or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE
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Calcium-binding.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 administration.",
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TISSUE SPECIFICITY: MAINLY PRESENT IN THE LIVER. WEAK EXPRESSION
WAS FOUND IN THE BRAIN AND LUNG, NOT FOUND IN THE KIDNEY.
DEVELOPMENTAL STAGE: PROTEIN AMOUNTS IN LIVER DECREASE
SIGNIFICANTLY WITH AGE.
SIMILARITY: BELONGS TO THE SMP-30 / CGR1 FAMILY.
SWISS-PROT entry
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FUNCTION: MAY PLAY A ROLE IN THE REGULATION OF ENZYMATIC ACIN THE LIVER. DECREASE OF SMP30 LEADS TO THE DYSREGULATION CALCIUM SIGNALING IN THE AGED LIVER.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                124 PAVLERHOGS
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                                                                                                                                                                                                                                                                                                                                                                                                          cuniculus (Rabbit).
Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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60.0%;
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Pred. No..
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Mismatches
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                                            CGR1 FAMILY.
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By similarity).
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tent is in no
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01-OCT-1993
01-NOV-1997
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RAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Shimokawa N., Yamaguchi M.;
Shimokawa N., Yamaguchi M.;
"Molecular cloning and sequencing of the cDNA coding
binding protein regucalcin from rat liver.";
FEBS Lett. 327:251-255(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                between the Swiss Institute of Bioinformatics and the EMBL outst
the European Bioinformatics Institute. There are no restrictions
use by non-profit institutions as long as its content is in
modified and this statement is not removed. Usage by and for com
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STRAIN-Wistar; TISSUE=Liver;
MEDLINE=93041931; PubMed=1420310;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-NOV-1997 (Rel.
15-DEC-1998 (Rel.
Senescence marker
                                                                                                                                                                                                                    Fujita T., Shirasawa T., Uchida K., Maruyama N., "Gene regulation of senescence marker protein-30 up regulation with tissue maturation and gradual
                                                                                                                                                                                                                                                                                                                                           Shimokawa N., Matsuda Y., Yamaguchi M.; "Genomic cloning and chromosomal assignment mol. Cell. Biochem. 151:157-163(1995).
                                                                                                                                                                                                                                                                                                                                                                                                       STRAIN=Sprague-Dawley; TISSUE=Kidney; MEDLINE=96150990; PubMed=8569761;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRAIN=Wistar; TISSUE=Liver; MEDLINE=93351639; PubMed=8348951;
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Mammalia; Eutheria; Rodentia;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EMBL; AB035445; BAA88079.1; -.
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                                                                                                                                                                                                                                                                                 MEDLINE=96386712; PubMed=8794449;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Fujita T., Shirasawa T., Uchida K., Maruyama N., "Isolation of cDNA clone encoding rat senescence (SMP30) and its tissue distribution.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NCBI_TaxID=10116;
                                                                                                                                                                                   Mech.
                                                                                                                                                                                                 with aging.";
                                                                                                                                                                                                                                                                                                            TISSUE SPECIFICITY.
                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE OF 117-299 FROM N.A.
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                                                                           h. Āgeing Dev. 87:219-229(1996).

FUNCTION: MAY PLAY A ROLE IN THE REGULATION OF ENZYMATIC AVENUE OF SMP30 LEADS TO THE DYSREGULATION OALCIUM SIGNALING IN THE AGED LIVER.

SUBCELLULAR LOCATION: Cytoplasmic.

TISSUE SPECIFICITY: HEPATOCYTES AND RENAL PROXIMAL TUBULAR
                      EPITHELIUM.
DEVELOPMENTAL STAGE
FOUND IN 5-DAY-OLD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OR SMP30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RAT
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      A PLATEU
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37, Last annotation update)
protein-30 (SMP-30) (Regucalcin)
                                           STAGE:
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    NEONATES.
                                         IN LIVER,
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Pred. No. 15;
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Sciurognathi; Muridae:
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ER, THE FIRST PEAK OF EXPRESSION WAS EXPRESSION INCREASES FROM DAY 7 AND 3-6.5 MOTH-OLD ADULTS EXPRESS ABOUT
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Muridae; Murinae; Rat
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Best Local S
Matches 6
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EMBL; D31662;
PIR; S27203; S
                                                                                 EMBL;
                                                                                                                              This SWISS-PROT entry is copyright. It is produced through a between the Swiss Institute of Bioinformatics and the EMBI the European Bioinformatics Institute. There are no restrict use by non-profit institutions as long as its content is modified and this statement is not removed. Usage by and for the statement is not removed.
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15-DEC-1998
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HOXO_/
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                                                                                                                     entities
                                                                                                                                                                                                                                                                            STRAIN=H16 / DSM 428 / ATCC 17699;
MEDLINE=93015670; PubMed=1383192;
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01-JUL-1993
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                                               SEQUENCE
                                                                                                          or send
                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                     Bacteria;
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Plasmid megaplasmid pHG1.
                                                                                                                                                                                                                                                                                                                                                                                                   Hydrogenase
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                                                          Plasmid
                                                                       PIR; G43255;
                                                                                                                                                                                                        - ! - SIMILARITY: BELONGS TO THE
                                                                                                                                                                                                                    Alcaligenes eutrophus H16.";
J. Bacteriol. 174:6277-6289(1992)
                                                                                                                                                                                                                                                       Kortlueke C.,
Friedrich B.;
                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=510
                                                                                                                                                                                                                                          gene complex coding
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SIMILARITY: BELONGS TO THE SMP-30 / CGR1 FAMILY.
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                                                                                  M96433;
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                                                                                                          s requires a license agreement (S an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                     megaplasmid pHG1.
a; Proteobacteria;
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BAA06507.1;
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                                                                                  AAA16467.1;
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26, Last sequence up
37, Last annotation
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33389
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            59.6%;
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60.0%;
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                                                                                                                                                                                                                                                                                                                                                     beta subdivision; Ralstonia group;
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Score 34; DB
Pred. No. 12;
3; Mismatches
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E3CF6D3FFCAE4E98
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RESULT 15
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DT 15-
Search completed: June 27, 2003, 17:17:08 Job time: 9.7 secs
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CC paportunistic pathogen.", Nicotinate D-ribosucleotide + diphosphate - nicotinate + 5-phospho-alpha-D-ribose 1-diphosphate - nicotinate + 5-phospho-alpha-D-ribose 1-diphosphate - STATHWAY: NAD biosynthesis, nicotinamide to NaMN; second step. CC -- SIMILARITY: BELONGS TO THE NAPRTASE FAMILY.
                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 6; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                    Pyridine nucleotide biosynthesis; Transferase; Glycosyltransferase; Complete proteome.
SEQUENCE 399 AA; 46095 MW; 49B2266CF47346DE CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EMBL; AE004905; AAG08304.1;
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Q9HUP4;
15-JUN-2002 (Rel. 41, Created)
15-JUN-2002 (Rel. 41, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Nicotinate phosphoribosyltransferase 1 (EC 2.4.2.11) (NAPRTase 1).
PNCB1 OR PA4919.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRAIN=ATCC 15692 / PAO1;
MEDLINE=20437337; PubMed=10984043;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
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                                                                                                                                                                                                       1 GPRVVERHOSA 11
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                                                                                                                                                                                                                                                                                          59.6%; Score 34; DB 1; Length 399; 54.5%; Pred. No. 32;
                                                                                                                                                                                                                                                                   Mismatches .
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Post-processing: Minimum Match 0%
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seq length:
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57
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SPTREMBL_21:*
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                                                                                                                                                                                                                                                                                                                                                                                                                                    sp_archea:*
sp_bacteria:*
sp_fungi:*
                                                                                                                                                                                                                                                                              sp_unclassified:*
sp_rvirus:*
sp_bacteriap:*
sp_archeap:*
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sp_virus:*
sp_vertebrate:*
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sp_phage:*
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7 Q8TXJ9
Q95VY1
                                                                                   Q8WW76
Q99K47
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 Q9vg74 drosophila
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Q95vy1 drosophila
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Q99k47 mus musculu
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Q919n1 brachydanio
Q91del arabidopsis
Q922k2 rattus norv
Q99199 mus musculu
Q8x454 cavia porce
Q9fc35 streptomyce
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                                           Q99K47 PRELIMINARY; PRT; 557 AA.
Q99K47;
01-JUN-2001 (TrEMBLrel. 17, Created)
01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Fibrinogen A alpha polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TISSUE=LIVER; Strausberg R.; Strausberg R.; Submitted (JAN-2002) to the EMBL/GenBank/DDBJ databases. EMBL; BC020764; AAH20764.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa;
Mammalia; Eutheria;
   Mus musculus (Mouse).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE
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01-OCT-2000
01-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Brachydanio rerio (Zebrafish)
Eukaryota; Metazoa; Chordata;
Actinopterygii; Neopterygii; T
Cyprinidae; Danio
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Dapla.
DAPlA.
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                                                                                                             eurosids II; Bra
NCBI_TaxID=3702;
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Mammalia; Eutheria;
           Stracke
                              STRAIN=CV. COLUMBIA;
                                                                                                                                                                        Eukaryota; Viridiplantae; Stre Spermatophyta; Magnoliophyta;
                                                                                                                                                                                                                              MYB108 OR F5E6.18.
Arabidopsis thaliana
                                                                                                                                                                                                                                                                                                                                      01-OCT-2000 (TrEMBLrel.
01-OCT-2000 (TrEMBLrel.
01-MAR-2002 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Q9LDE1
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MEDLINE=20373792; PubMed=10917738;
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[1]
                                                             SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                      putative).
                                                                                                                                                                                                                                                                                                             Putative transcription
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Inohara N., Nunez G.
"Genes with homology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 issp;
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L; AF231127; AAF
UENCE 105 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               _TaxID=7955;
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P02671; 1FZA.
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R., Weisshaar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity 7; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GPRVVERHQSA 11
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                                                                                                                                               Brassicales;
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AAF66957.1; -.
AA; 11367 MW; 715D
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Rodentia;
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                                                                                                                                                                   (Mouse-ear cress)
htae; Streptophyta; Embry
liophyta; eudicotyledons;
                                                                                                                                                                                                                                                                                                               factor
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    Created)
    Last sequence update)
    Last annotation update)
    MYB108 (Myb-related protein,

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                                                                                                                                               Brassicaceae;
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m; Craniata; Verte
; Teleostei; Ostar
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Last annotation updat
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Pred. No.
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0.046;
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Ostariophysi;
                                                                                                                                                                        Embryophyta; Tracheophyta;
edons; core eudicots; Rosid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB
                                                                                                                                            Arabidopsis
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Q9Z2T2;
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                                                                                                                                                                                                                                            InterPro; IPR003280; K+channel_2pore
InterPro; IPR001522; K+channel_pore.
InterPro; IPR001632; K+channel_nlg.
InterPro; IPR001779; TWIK1_channel.
Pfam; PF00520; ion_trans; I.
PRINTS; PR01333; 2FOREKCHANEL.
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SMART; SM0395; SANT; 2.
PROSITE; PS00037; MYB_1; UNKNOWN
PROSITE; PS00334; MYB_2; 1.
PROSITE; PS50090; MYB_3; 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lin X., Kaul S., Town C.D., Benito M., Creasy T.H., Haas B.J., Maiti R., Ronning C.M., Koo H., Fujii C.Y., Utterback T.R., Barnstead M.E., Bowman C.L., White O., Nierman W.C., Fraser C.M. "Arabidopsis thaliana chromosome 3 BAC F586 genomic sequence."; Submitted (JAN-2001) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                      PRINTS; PR01333; PRINTS; PR01096; PRINTS; PR01587;
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Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Putative potassium channel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-MAY-1999 (TrEMBLrel. 10,
01-MAY-1999 (TrEMBLrel. 10,
01-JUN-2002 (TrEMBLrel. 21,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HSSB; PO1103; 1POM.
HSSB; PO1103; 1POM.
InterPro; IPR001005; Myb DNA binding.
InterPro; IPR001005; Myb DNA-binding; 2.
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                                                                                                                                      SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  !- SUBCELLULAR LOCATION: NUCLEAR (! !- SIMILARITY: CONTAINS 2 MYB-LIKE
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RVVERHQSA 11
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77.8%;
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Pred. No. 39;
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InterPro; IPR001522; K+channel pore.
InterPro; IPR001622; K+channel nlg.
InterPro; IPR000636; M+channel nlg.
InterPro; IPR001779; TWIK1 channel.
Pfam; PF00520; ion trans; I
PRINTS; PR01333; 2POREKCHANEL.
PRINTS; PR01333; 2POREKCHANNEL.
PRINTS; PR01586; TWIKICHANNEL.
PRINTS; PR01586; TWIKICHANNEL.
                                                                                                                                                                                                                                                                               Q8R454;
01-JUN-2002
01-JUN-2002
01-JUN-2002
  Q9FC35
                                                                                                                                                  EMBL;
                                                                                                                                                        SEQUENCE FROM N.A.

Derst C., Rajan S., Preisig-Mueller R.;
"Cloning and sequencing of guinea pig TWIK channels.";
"Cloning and sequencing of guinea pig TWIK channels.";
Submitted (JAN-2002) to the EMBL/GenBank/DDBJ database
                                                                                                                                                                                                                         Cavia porcellus (Guinea pig).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
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Mammalia; Eutheria; Rodentia;
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01-JUN-2001
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InterPro; IPR003280; K+
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NCE 336 AA; 38102 M
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Hystricognathi; Caviidae; Cavia.
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Nature 417:141-147(2002).
EMBL; AL391039; CAC01546.1; -.
InterPro, IPR001225; FA_desaturase.
Pfam; PF00487; FA_desaturase; 1.
ProDom; PD001081; FA_desaturase; 1.
ProDom; PD001081; FA_desaturase; 1.
SEQUENCE 347 AA; 37990 MW; 201E2D
                                                                                            01-JUN-2001
01-JUN-2001
01-JUN-2001
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Q9C6A4;
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01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation updat
Putative delta fatty acid desaturase.
$CO7048 OR $C4G1.14C.
$treptomyces coelicolor.
          Arabidopsis thaliana (Mouse-ear cress).

Bukaryota; Viridiplantae; Streptophyta; Embryophyta;

Spermatophyta; Magnoliophyta; eudicotyledons; core eu
eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bentley S.D., Chater K.F., Cerdeno-Tarraga A.-M., Challis G.L., Thomson N.R., James K.D., Harris D.E., Quali M.A., Kieser H., Harris D.E., Quali M.A., Kieser H., Collins M. Harper D., Bateman A., Brown S., Chandra G., Chen C.W., Collins M. Cronin A., Fraser A., Goble A., Hidalgo J., Hornsby T., Howarth S. Huang C.-H., Kieser T., Larke L., Murphy L., Oliver K., O'Neil S., Rabhinowirsch E. Batandram M. Butherford V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
STRAIN=A3(2) / M145;
Bentley S.D., Chater K.F.,
Thomson N.R., James K.D., I
                                                                  Hypothetical F9E11.8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=97000351; PubMed=8843436; Redenbach M., Kieser H.M., Denapaite
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Brown S.P.;
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NCBI_TaxID=1902;
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"A set of ordered cosmids and a detailed genetic and the 8 Mb Streptomyces coelicolor A3(2) chromosome.";

Mol. Microbiol. 21:77-96(1996).
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EMBL/GenBank/DDBJ databases.
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                                                                                                                        Created)
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                        eudicots;
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RESULT 11
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A MEDLINE-21016719; PubMed=11130712;

A White O., Alonso J., Altafi H., Araujo R., Bowana C.L., Brooks S.Y.,

A Buehler E., Chan A., Chao O., Chen H., Cheuk R.E., Chin C.W.,

A Chung M.K., Conn L., Conway A.B., Conway A.R., Creasy T.H., Dewar K.,

A Chung M.K., Conn L., Conway A.B., Conway A.R., Creasy T.H., Dewar K.,

A Gill J.E., Goldsmith A.D., Haas B., Hansen N.F., Hughes B., Huizar L.,

RA Gill J.E., Goldsmith A.D., Haas B., Hansen N.F., Hughes B., Huizar L.,

RA Kim C.J., Koo H.L., Kremenetskala I., Kurtz D.B., Kwan A., Lam B.,

RA Langin-Hooper S., Lee A., Lee J.M., Lenz C.A., Li J.H., Li Y.-P.,

RA Lin X., Liu S.X., Liu Z.A., Luros J.S., Maiti R., Marziali A.,

RA Militscher J., Miranda M., Nguyen M., Nierman W.C., Osborne B.I.,

RA Pai G., Peterson J., Pham P.K., Rizzo M., Rooney T., Rowley D.,

RA Pai G., Peterson J., Pham P.K., Rizzo M., Rooney T., Rowley B.I.,

RA Sun H., Tallon L.J., Tambunga G., Toriumi M.J., Town C.D.,

RA Utterback T., Van Aken S., Vayaberg M., Vysotekaia V.S., Walker M.,

"Sequence and analysis of chromosome 1 of the plant Arabidopsis
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01-DEC-2001
01-DEC-2001
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EMBL; X70431; CAA49881.1; -
CHAIN 2 65
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Q53379;
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SEQUENCE FROM N.A.
STRAIN=CV. COLUMBIA
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EMBL; AC079678; AAG51875.1;
InterPro; IPR001798; Kelch.
Pfam; PF01344; Kelch; 5.
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Pred. No. 18;
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Pred. No. 1e+02;
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RX MEDLINE=2108560; Pubmed=1127851;
RA Rawai J., Shinagawa A., Shibata K., Yoshino M., Itoh M., Ishii Y.,
RA Arakawa T., Hara A., Fukunishi Y., Konno H., Adachi J., Fukuda S.,
RA Aizawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I.,
RA Aizawa K., Izawa M., Nishi K., Kiyosawa H., Kasukawa T., Saito R.,
RA Saito T., Okazaki Y., Gojobori T., Bono H., Kasukawa T., Saito R.,
RA Kadota K., Matsuda H.A., Ashburner M., Batalov S., Casavant T.,
RA Kadota K., Matsuda H.A., Ashburner M., Batalov S., Cosavant T.,
RA Kuchl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Kuchl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Schriml I.M., Staubli F., Suzuki R., Tonita M., Wagner L., Washio T.,
RA Sakai K., Okido T., Furuno M., Aono H., Baldarelli R., Barsh G.,
RA Blake J., Boffelli D., Bojunga N., Carninci P., de Bonaldo M.F.,
RA Brownstein M.J., Bult C., Fletcher C., Fujita M., Gariboldi M.,
RA Lyons P., Marchionni L., Mashima J., Mazzarelli J., Mombaerts P.,
RA Sasaki H., Sato K., Schenbach C., Seya T., Shibata Y., Storch K.-F.,
RA Mynshaw-Boris A., Yoshida K., Hasegawa Y., Kawaji H., Kohtsuki S.,
RA Havashizaki Y.;
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Misawa H., Yamaguchi M.;
Wihe gene family encoding the c.
Submitted (FEB-2000) to the EMB
EMBL; AB037934; BAA90692.1; -
SEQUENCE 299 AA; 33390 MW;
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Q925W3;
Q1-DEC-2001
01-DEC-2001
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Mammalia; Eutheria;
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Pfam; PF01476; LysM; 1.
SMART; SM00257; LysM; 1.
SEQUENCE 215 AA; 23765
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01-JUN-2001
01-JUN-2001
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Mammalia; Eutheria;
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                Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (TrEMBLrel. (TrEMBLrel. (TrEMBLrel.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               N.A.
6J; TISSUE=STOMACH;
                                                                                                                                                                                                                                                                                                                                                                                  (Rat).
                                                                                                                                                                                                                                                                                                                             Chordata;
Rodentia;
                                 60.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     23765 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Chordata;
Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       61.4%;
75.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                               19,
19,
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17,
17,
  Score 35; DB Pred. No. 86; 2; Mismatches
                                                                                                                                                            he calcium-binding
EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                            Last sequence update)
Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1;
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Last sequence
Last anno
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 35; DB Pred. No. 61;
                                                                                                                                                                                                                                                                                                               Craniata; Vertebrata; Euteleostomi;
Sciurognathi; Muridae; Murinae; Rattus.
                                                                                                         5D8F2D95FCA4EE35 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DC011B368AD1B976 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sequence update) annotation updat
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                                                                                                                                                            protein regucalcin.";
databases.
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                                                    Length 299;
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,
Gaps
0;
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N-terminal

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RA Adams N.D., Celniker S.E., Holt R.A., Evans C.A., Gocayne J.D.,
RA Adams N.D., Celniker S.E., Holt R.A., Evans C.A., Gocayne J.D.,
RA Adams N.D., Celniker S.E., Holt R.A., Evans C.A., Golale R.F.,
RA Adams N.D., Celniker S.E., Holt R.A., Evans C.A., Galle R.F.,
RA George R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N.,
RA Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X.,
RA Burton R.C., Rogers Y.-H.C., Blazej R.G., Champe M., Pfeiffer B.D.,
RA Barlandon R.C., Rogers Y.-H.C., Blazej R.G., Champe M., Pfeiffer B.D.,
RA Barlandon R.C., Rogers Y.-H.C., Blazej R.G., Champe M., Pfeiffer B.D.,
RA Barlandon R.C., Rogers Y.-H.C., Blazej R.G., Champe M., Pfeiffer B.D.,
RA Barlandon R.C., Rogers Y.-H.C., Blazej R.G., Champe M., Pfeiffer B.D.,
RA Barlandon R.D., Barnandal J., Bayraktaroglu L., Beasley E.M.,
RA Barlaw R.M., Basu A.N., Baxendale J., Bayraktaroglu L., Beasley E.M.,
RA Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S.,
RA Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P.,
RA Berson K.Y., Besons P., W. Berman B.P., Bhandari D., Bolshakov S.,
RA Burtis K.C., Busam D.A., Burler H., Cadleu E., Center A., Chandra I.,
RA de Pablos B., Delcher A., Deng Z., Mays A.D., Dew I., Dietz S.M.,
RA de Pablos B., Delcher A., Demes M., Dugan-Rocha S., Dunkov B.C., Dunn P.,
RA de Pablos K., Doup L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P.,
RA Glodek A., Gong F., Gorrell J.H., Gu Z., Glann P., Harris M.,
RA Harris N.L., Harvey D., Heiman T.J., Hernandez J.R., Houck J.,
RA Harris N.L., Katuston K.A., Howland T.J., Wei M.-H., Diegwam C.,
RA Harris N.L., McIntosh T.C., McLeed M., H., McBherson D.,
RA Kimmel B.E., Kodira C.D., Kraft C., Kraviz S., Kulp D., Lai Z.,
RA Lasko P., Lei Y., Levitsky A.A., Li J., Li Z., Liang Y., Lin X.,
RA Harris N.L., McIntosh T.C., McLeed M., McBeerson D.L.,
RA Harris N., McMarty D.M., McHerson D.L.,
RA McHulov G., Milshina N.V., Mobarry L., Moshrefi A.,
RA McHulov G., Milshina N.V., Mobarry L., Mushry D.M., Nelson D.L.,
RA Mans R., Weissen M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 13
Q9VG74
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                            Query Match
Best Local :
     Matches
                                                                                                               Pfam; PF01354; Antifreeze; 1.
Pfam; PF03102; NeuB; 1.
SEQUENCE 338 AA; 37535 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Drosophila melanogaster (Fruit fly).
Bukaryota, Metazoa; Arthropoda, Tracheata; He
Pterygota, Neoptera, Endopterygota, Diptera;
Ephydroidea; Drosophilidae; Drosophila.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CG5232 protein.
NEU5AC OR CG5232.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-MAY-2000 (TrEMBLrel.
01-JUN-2002 (TrEMBLrel.
                                                                                                                                                                                                                                                FlyBase; FBgn0038045; Neu5Ac.
                                                                                                                                                                                                                                                                            EMBL; AE003695; AAF54811.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=20196006;
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                                                                                                                                                                                            InterPro; IPR000029; AntifreezeIII.
InterPro; IPR004144; NeuB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             N
  Similarity 6; Conserv
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     Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PubMed=10731132;
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                                                                                                               37535 MW;
                      61.4%;
75.0%;
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13,
21,
Score 35; DB Pred. No. 97; Mismatches
  ۲.
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Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Created)
                                                                                                            95F03C4FE42956C2 CRC64;
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a; Brachycera; Muscomorpha;
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  1;
                                                   Length 338;
  Indels
0;
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Q95VY1
Q95VY2
Q95VY2
ID Q95VY2
AC Q95VY2
AC Q95VY2
AC Q95VY2
DT 01-DE
DT 01-DE
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ERN (
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Q8TXJ9
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A
STRAIN=OREGON R;
Kim K., Lawrence (
Palter K.B.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Q95VY1;
Q95VY1;
01-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
SEQUENCE FROM N.A.
STRAIN=AV19 / DSM 6324 / JCM 9639;
STRAIN=AV19 / DSM 6324 / JCM 9639;
STRAIN=21927647; PubMed=11930014;
MEDLINE=21927647; PubMed=11930014;
Slebarev A.I., Mezhevaya K.V., Makarova K.S., Polushin N.N.,
Slebarev A.I., Mezhevaya K.V., Belova G.I., Aravind L.,
Shcherbinina O.V., Shakhova V.V., Belova G.I., Aravind L.,
Shcherbinina O.V., Shakhova V.V., Belova G.I., Aravind L.,
Shcherbinina O.V., Shakhova V.V., Belova G.I., Stetter
                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_T
                      InterPro; IPR000029; AntifreezeIII.
InterPro; IPR004144; NeuB.
Pfam; PF01354; Antifreeze; 1.
                                                                                                                             "Expression of a functional Drosophila acid (NeuSAc) phosphate synthase gene: acid biosynthetic ability in insects."; Glycobiology 0:0-0(2001).
EMBL; AF397531; AAK92125.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;
Pterygota; Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Neu5Ac synthase.
NEU5AC OR CG5232.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-DEC-2001 (TrEMBLrel. 19, 01-DEC-2001 (TrEMBLrel. 19, 01-MAR-2002 (TrEMBLrel. 20,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Methanopyrus kandleri.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RIO1-like serine/threonine DNA-binding HTH domain.
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01-JUN-2002 (TrEMBLrel. 21, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 QBTXJ9;
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                                                                                                         FlyBase; FBgn0038045; NeuSAc
                                                                                                                                                                                                                                                                                                                                                                                                                                             Ephydroidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Drosophila melanogaster (Fruit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=2320;
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                                                                                                                                                                                                                                                                                                                                                                                                                TaxID=7227;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       194
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GPRVVERH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                       Drosophilidae; Drosophila
                                                                                                                                                                                                                                                                                                                                R.A.
                                                                                                                                                                                                                                                                                                   S.M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       201
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54.5%;
                                                                                                                                                                                                                                                                                                   Park J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       protein kinase fused to an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Last sequence update)
Last annotation update)
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1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             fly)
                                                                                                                                                                                                                                                                                                   Pitts L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                35; DB 17;
No. 1.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         372
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Methanopyrales; Methanopyraceae;
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Evidence for
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                                                                                                                                                                                                                                                                                                   Vann W.F.,
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Length 368 Indels

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Gaps

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N-acetylneuraminic endogenous sialic Betenbaugh M.J.,

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Search completed: June 27, 2003, 17:18:03 Job time : 28.4 secs	1 GPRVVERH 8 : 228 GARIVERH 235	Query Match 61.4%; Score 35; DB 5; Length 372; Best Local Similarity 75.0%; Pred. No. 1.1e+02; Matches 6; Conservative 1; Mismatches 1; Indels	SEQUENCE
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                                 Gapop 10.0 , Gapext 0.5
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57
908470 segs, 133250620 residues
                                                                                          1 GPRVVERHQSA 11
                                                                                                                                                                                                                                                                                          GenCore version 5.1.6
Copyright (c) 1993 - 2003 Compugen Ltd
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Minimum DB seq length: 0
Maximum DB seq length: 200000000 Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 s summaries

Total number of hits satisfying chosen parameters:

908470

Database : 11: 12: 13: 14: 15: 16: 17: 19: A_Geneseq_101002:*

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and is derived Pred. No. is the number of results predicted by chance to have a ster than or equal to the score of the result being printed, rived by analysis of the total score distribution.

SUMMARIES

10	ø	8	7	σ	ຫ	4.	ω	N		Result No.
57	57	57	57	57	57	57	57	57	57	Score
100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	Query
643	360	78	78	23	15	្រី	ŭ,	ঠ		Query Match Length DB
15	21	23	23	21	21	14	21	19	22	BB
AAR60020	AAB54135	AA014005	AA021122	AAY54925	AAY95518	AAR69318	AAB13511	AAW50593	ABB76978	Ib
Fibronectin. Homo	Human pancreatic c	Human fibrinogen E	Fibrinogen E alpha	Peptide ligand for	Fibrin polymerizat	Gp IIb/IIIa recept	Fibrin derived pep	GPIIb/IIIa recepto	Residues 36-46 of	Description .

45	44	43	42	41	40	39	38	37	36	35		33	32	31	30		28	27	26	25	24	23	22	21	20	19	18	17	16	15	14 .	13	12	11
ω 5	3 5	35	ω	35	36	ω 6	36	36	36	36	36	37	37	37	37	37	38	38	38	38	ა 8	38	ω 8	38	41	44	44	44	44	44	44	53	57	57
61.4	61.4	61.4	61.4	۲.	ω	Ψ	ω.	w.	ω.	63.2	ü	٠.	4.	٠.	4.		66.7		66.7		66.7			•	71.9		٠	•	77.2	•	•	٠	100.0	100.0
214	131	107	74	73	132	50	24	12	œ	7	7	336	336	325	323	140	299	299	299	24	24	24	24	24	10	24	24	24	24	24	17	10		644
21	22	23	22	22	22	22				11	œ	21	18	22	23	22	20	18	16	23	23	23	23	23	19	23	23	23	23	23	23	16	21	1.6
AAY99923	AAU55722	ABP32045	ABG26011	ABG26014	ABG26016	ABG28568	AAW64264	AAP71099	AAR65794	AAR06526	AAP71315	AAY79673	AAW23397	ABG20112	AAU93178	ABG04700	AAY06995	AAW14475	AAR74219	AA021121	AAO21120	AA021115	AAO21114	AA021113	AAW65536	AA021119	AA021118	AA021117	AAO21116	AA021112	AAE19125	AAW11932	AAY82891	AAR82244
Peptide encoded by	Propionibacterium	Human RNA polymera	Novel human diagno	Novel human diagno	^	Novel human diagno	Murine 34-, 40 and	Functional site re	Fibrin-specific ep	Peptide antigenic	Sequence of fibrin	Human potassium ch	TWIK-1 potassium c	Novel human diagno		Novel human diagno		Human senility mar	Human ageing marke	Anti-angiogenic al	Anti-angiogenic al	Anti-angiogenic al	a	Anti-angiogenic al	Biotinyl-fibrin-an	Anti-angiogenic al	Thrombin peptide.	oge	AlphaE subunit of	Human fibrinogen A				

ALIGNMENTS

RESULT 1 ABB76978 ABB76978; ABB76978 standard; peptide; 11 AA

Residues 36-46 of human fibrinogen alpha chain precursor

22-JUL-2002 (first entry)

Anti-arthritic; anti-inflammatory; fibrin; rheumatoid polyarthritis; human; fibrinogen alpha chain.

Homo sapiens.

01-JUL-1999; 01-JUL-1999; 05-JAN-2001 FR2795735-A1 99FR-0008470 99FR-0008470.

WPI; 2001-114394/13. Serre G, Sebbag M;

(UYTO-) UNIV TOULOUSE SABATIER PAUL.

New citrulline-containing polypeptide from fibrin, useful for diagnosis and treatment of rheumatoid polyarthritis

Example 1; Page 11; 23pp; French.

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RESULT 2
AAW50593
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Best Local S
Matches 11
                                                                                                                                                                                                                           07-JUN-1995;

08-FEB-1991;

27-NOV-1991;

21-MAY-1992;

22-JUN-1994;

11-JUL-1994;

07-JUN-1995;
                                                                                                     Reagent use
peptide cov
technetium
The invention relates to a reagent for preparing a thrombus imaging agent. It comprises: (a) a specific binding peptide (BB) having a sequence of 4-100 amino acids; and (b) a technetium-99m binding moiety (TBM) covalently linked to the peptide which binds to a component of a thrombus. TBM forms an electrically neutral complex with technetium-99m
                                                                                                                                                                                                                                                                                                                                                                   US573,6122 A.
                                                                                                                                                     WPI; 1998-239148/21.
                                                                               Example
                                                                                                                                                                               Dean RT,
                                                                                                                                                                                                       (DIAT-)
                                                                                                                                                                                                                                                                                                                             07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                 07-APR-1998.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GPIIb/IIIa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to a citrulline (Cit) containing polypeptide derived from all or part of the alpha- or beta-chains of fibrin by substitution of at least one arginine residue by Cit. The Cit containing polypeptides can be used for in vitro diagnosis of rheumatoit polyarthritis (RP), by detecting disease-specific autoantibodies, and therapeutically for neutralising the RP-associated autoimmune response. The present sequence is a fragment (residues 3636) of the human fibrinogen alpha chain precursor, which was used in an example from the invention
                                                                                                                                                                                                                                                                                                                                                                                                                              Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Technetium-99M label;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16-JUL-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAW50593;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAW50593 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                useful for preparing thrombus imaging agent - covalently linked to technetium binding moiety ium labelled and binds to thrombus component
                                                                               2;
                                                                                                                                                                                                     DIATIDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        moiety;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GPRVVERHOSA 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GPRVVERHQSA
                                                                                                                                                                             Lister-James
                                                                           Columns 11-12; 18pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                           95US-0482880.
91US-0653012.
91US-0807062.
92US-0886052.
94US-0264176.
94US-0273274.
95US-0480551.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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                                                                                                                                                                                                                                                                                                                             95US-0482880
                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
12
/note= "Side chain"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         branched
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ligand with technetium-99M binding moiety.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 thrombus imaging;
                                                                                                                                                                            ٦
                                                                                                                                                                                                                                                                                                                                                                                              "Side chain of Lys is linked to an identical peptide chain to form a branched peptide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 57;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ;GPIIb/IIIA receptor; ligand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.00014;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               rheumatoid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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GPRVVERHOSA

11

Query Match Best Local S Matches 11

l Similarity

Conservative

0;

Mismatches

100.

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Score 57; I Pred. No. 0.

DB 21;).00017;

Length 13;

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Gaps

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RESULT 3
AAB13511
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
Matches 11
                           The present sequence is a peptide derivative of fibrin. It was used as a example of the method of the invention, which involves the production of compositions comprising a polypeptide with a water soluble polymer covalently bound to the N-terminal alpha carbon by a hydrazone or an oxime (or the reduced version of each) bond. This is useful as previous methods were more time consuming and labour-intensive, as well as being less precise. The composition can be used to produce diagnostic and
Sequence
                                                                                                                                                        Covalently binding a water-soluble polymer to the N-terminal alpha-carbon atom of a polypeptide for producing therapeutic and diagnostic agents, by contacting the transaminated polypeptide with the polymer to form a hydrazone bond -
                       therapeutic
                                                                                                                                       Example
                                                                                                                                                                                                                                                                                                      04-AUG-1997;
                                                                                                                                                                                                                                                  Wei Z,
                                                                                                                                                                                                                                                                                                                              04-AUG-1997;
                                                                                                                                                                                                                                                                       (ORTH ) ORTHOPMCNEIL PHARM INC
                                                                                                                                                                                                                                                                                                                                                       20-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                             US6077939-A
                                                                                                                                                                                                                                                                                                                                                                                                                                Cross-links
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Fibrin derivative; diagnostic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Fibrin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              02-NOV-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAB13511 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The small size of the peptide component of the reagent enhances blood and background tissue clearance and it is easy to prepare. This sequence represents a GPIIb/IIIa receptor binding peptide covalently linked to a technetium-99m binding moiety.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAB13511;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    derived
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11;
                                                                                                                                 7; Column 14; 21pp; English.
                                                                                                                                                                                                                                                  Ghosh-Dastidar P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GPRVVERHQSA
13
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                      agents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
ΑA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                  97US-0905310
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                                                                                                                                                                                                                                                                                                                                                                                                    /note= "this peptide is
peptide by a disulphide
                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%;
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                                                                                                                                                                                                                                                 Menon-Rudolph
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Pred. No. 0.0
); Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          agent; therapeutic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 19;
0.00016;
                                                                                                                                                                                                                                                                                                                                                                                                  linked to a second copy bond to form a dimer"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           reagent enhances blood
to prepare. This
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length
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RESULT 5
AAY95518
ID AAY9
XX
AC AAY9
                                                                                                                              RESULT 4
AAR69318
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                                                                                                    Query Match
Best Local
                                                                                            Matches
                                                                                                                                                                                                                                              Reagent for scintigraphic imaging of thrombi with 99m technetium - comprises synthetic peptide which binds to thrombus, covalently coupled to metal binding gp.; rapidly cleared from blood and
                                                                                                                                                   The invention relates to reagents for scintigraphic imaging of a thrombus in-vivo, comprising (A) a specific binding compound capable binding to at least one component of a thrombus, covalently linked to (B) a technetium-99m-binding moiety. Specific peptides constituting the reagents are claimed as new. The present peptide is one such peptide, in which the Cys (Acm) - Gly-Cys (Acm) moiety is the 99m-Tc binding moiety and the residue constitutes the thrombus-
                  AAY95518
                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                 WPI; 1993-386229/48
                                                                                                                                                                                                                                                                                                                                                    21-MAY-1993;
                                                                                                                                                                                                                                                                                                                                                                                                              Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 technetium-99m;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Scintigraphy;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAR69318;
                                                                                                                                             binding component
                                                                                                                                                                                                                      Claim 19; Page 42; 61pp; English.
                                                                                                                                                                                                                                                                                                                                   21-MAY-1992;
                                                                                                                                                                                                                                                                                                                                                                    25-NOV-1993.
                                                                                                                                                                                                                                                                                                                                                                                     WO9323085-A
                                                                                                                                                                                                                                                                                                                                                                                                                               Modified-site
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                                                                                                                                                                                                                                                                                                                    (DIAT-) DIATECH
                                                                            .
1
                                                                                           l Similarity
                  standard;
                                                                           GPRVVERHQSA 11
                                                           GPRVVERHQSA 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GPRVVERHQSA
                                                                                                                              15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       receptor ligand used in scintigraphic imaging of thrombi
                                                                                           Conservative
                                                                                                                              A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              thrombus; thrombi; imaging; s
m; radiolabelled; Gp IIb/IIIa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                    92US-0886752
                                                                                                                                                                                                                                                                                                                                                    93WO-US04794.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                /note=
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                 Peptide; 15
                                                                                                   100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                    "S-acetamidomethyl-Cys-amide"
                                                                                                                                                                                                                                                                                                                                                                                                                                     "the omega-amino of this the peptide GPRVVERHQSA"
                                                                                                                                                                                                                                                                                                                                                                                                                     "S-acetamidomethyl-Cys"
                                                                                           ,
,
                                                                                                   Score 57; I
Pred. No. 0.
                                                                                           Mismatches
                                                                                                  DB 14;
0.0002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              specific binding;
a receptor ligand.
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                                                                                                          Length 15;
                                                                                           Indels
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                                                                                                                                                                                                                                                        covalently
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                                                                                          Gaps
                                                                                                                                                                                              of.
                                                                                          0
RESULT 6
AAY54925
ID AAY5
XX
AC AAY5
AC AAY5
XX
DT 15-F
XX
DE Pept
                                                                                          밁
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                                                                                                                                                              Matches
                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                   21-MAY-1992;
21-MAY-1993;
05-JAN-1995;
                                                                                                                                                                                                                                                                                                                  WPI;
Peptide ligand for fibrin polymerisation site.
                                 AAY54925;
                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                  Lister-James J,
                                                                                                                                                                                                                                                                                                                                                                                                                     04-JUL-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                      US6083481-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 embolism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10-OCT-2000
               15-FEB-2000
                                                 AAY54925 standard;
                                                                                                                                                                                                                                                                                                                                                   (DIAT-) DIATIDE
                                                                                                                                                                                                                                                                                                                                                                                                    27-AUG-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Thrombus;
                                                                                                                                    Local
                                                                                                                                                                                                                                                                                                                  2000-498061/44.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         polymerization
                                                                                                                                   Similarity
                                                                                          GPRVVERHQSA 11
                                                                                                         GPRVVERHOSA
                                                                                                                                                             15 AA;
                                                                                                                                                                                                                                                                                                                                                    INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note=
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The present sequence is that of a ligand for the fibrin polymerization site. The invention relates to radiolabeled reagents that are scintigraphic imaging agents for imaging sites of thrombus formation in vivo. The reagents each comprise a specific binding compound, such as the present peptide, that is capable of binding to at least 1 component of a thrombus, and which is covalently linked to a radiolabel-binding moiety. A method for using such a reagent labeled with technetium-99m to image a thrombus site in a mammalian body is claimed.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Composition comprises technetium-99m and polyamide reagent which bir to thrombi, useful as scintigraphic imaging agent for imaging sites thrombus formation in vivo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Column 10; 27pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      imaging; radioimaging; thrombosis; thromboembolism;
diagnosis; technetium-99m; fibrin ligand.
                                                                                                                                                                                                                                                                                                           Conservative
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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93WO-US04794.
95US-0335832.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                              peptide;
                                                                                                                                                                                                                                                                                                         100.0%; P
                                                                                                                                                                                                                                                                                                                                                100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "2 copies of attached to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Cys (Acm) "
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                                                                                              23
                                                                                                                                                                                                                                                                                                              Score 57; DB 21; Pred. No. 0.0002; Mismatches 0
                                                                                                A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       this peptide are covalently the Lys residue at position
                                                                                                                                                                                                                                                                                                                                                             DB 21;
                                                                                                                                                                                                                                                                                                                                                        Length
                                                                                                                                                                                                                                                                                                                Indels
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                                                                                                                                                                                                                                                                                                                Gaps
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RESULT 7
AAO21122
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This sequence represents a peptide ligand for the fibrin polymerisation site. The invention relates to a complex (A) for thrombus imaging comprises technetium-99m (TC-99m) completed with a reagent comprising a peptide (P) with 4 to 100 amino acids sequence and a TC-99m binding moiety covalently bound to (P). (P) is selected from a linear peptide ligand for a GPIIb/IIIa receptor not comprising the amino acid sequence (arginine-glycine-aspartate), a peptide ligand for a polymerisation site of fibrin, and a cyclic peptide ligand for the GPIIb/IIIa receptor. The thrombus imaging reagents provided by the present invention can be used for visualising thrombi in a mammalian body when TC-99m is labelled.
01-SEP-2000; 2000GB-0021475
                           03-SEP-2001; 2001WO-GB03926.
                                                                                                                                        gene
                                                                                                                                                                                                               Fibrinogen E alpha-chain protein sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A complex used for thrombus imaging comprises technetium-99m complexed with a peptide ligand for \mbox{GPIIb/IIIa} receptor -
                                                                                                              Unidentified
                                                                                                                                                                                          Cytostatic;
                                                                                                                                                                                                                                                                                               AAO21122 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2000-021733/02.
                                                           07-MAR-2002
                                                                                    WO200218440-A1
                                                                                                                                                   neovascular glaucoma;
                                                                                                                                                                            antipsoriatic; antidiabetic;
                                                                                                                                                                                                                                            19-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11-JUL-1994;
21-MAY-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19-OCT-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Thrombus imaging; fibrin polymerisation site; technetium-99m; Tc-99m; GPIIb/IIIa receptor; cyclic peptide ligand.
                                                                                                                                                            fibrinogen; cancer; angiogenesis; chemotherapeutic agent;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (DIAT-) DIATIDE INC
                                                                                                                                    therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       8
                                                                                                                                                                                                                                                                                                                                                                                                                11;
                                                                                                                                                                                                                                                                                                                                                                                      1 GPRVVERHOSA 11
                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                            GPRVVERHOSA 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Lister-James
                                                                                                                                                         antitumour; ophthalmological; antiarthritic; antirneumac.ic: antidiabetic; anti-angiogenic; alphal-24 peptide; tumc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23
                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; ilarity 100.0%; Conservative (
                                                                                                                                                                                                                                           (first entry)
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92US-0886052.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       95US-0484773.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   32; 18pp; English
                                                                                                                                                   rheumatoid
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                                                                                                                                                                                                                                                                                               78
                                                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                                                           Score 57;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                arthritis; diabetic retinopathy;
                                                                                                                                                                                                                                                                                                                                                                                                                           0.00032;
                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 23;
                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                      antirheumatic;
                                                                                                                                                                                                                                                                                                                                                                                                            0,
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RESULT 8
AAO14005
PF XXX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CC alpha 1.24 peptide sequence derived from fibrinogen. The peptide of comprising ADSGEGDFLAEGGGVRGPRVVERH (alpha 1.24 peptide) is useful for CC manufacturing a medicament for treating cancer. The peptide, its encoding CC DNA, a pharmaceutical composition and vector of the invention are useful CC angiogenesis, where the therapeutic agents are administered to the animal CC angiogenesis, where the therapeutic agents are administered to the animal CC angiogenesis, where the therapeutic agents are administered to the animal CC angiogenesis, where the therapeutic agent are conjugated, associated CC or crosslinked to a chemotherapeutic agent. The method is preferably CC carried out for inhibiting tumour angiogenesis in tumours. The peptide, its encoding DNA, a pharmaceutical composition and vector of the invention are useful for inhibiting angiogenesis in conditions of C neovascular glaucoma, rheumatoid arthritis, psoriasis, or diabetic C retinopathy. The DNA of the alpha 1.24 peptide can be used in gene C therapy. This sequence represents a fibrinogen E alpha-chain protein C sequence of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local S
Matches 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 200
N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   09-NOV-2000;
27-NOV-2000;
23-JUL-2001;
                                                                                                                                  glomerulonephritis.
                                                                                                                                                  ovarian hyperstimulation; obesity; hepatitis; pneumonia
                                                                                                                                                                                     neovascular glaucoma; diabetic
                                                                                                                                                                                                                                                          Human fibrinogen E-fragment; alpha-chain; anti-angiogenesis;
                                                                                                                                                                                                                                                                                                                                 28-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                      age-related macular degeneration;
                                                                                                                                                                                                       tumour identification;
                                                                                                                                                                                                                       thrombin; breast carcinoma; cancer;
                                                                                                                                                                                                                                        angiogenesis; blood vessel development; cellular matrix;
                                                                                                                                                                                                                                                                                            Human fibrinogen E-fragment alpha-chain amino acids 1-78
                                                                                                                                                                                                                                                                                                                                                                                              AAO14005 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel polypeptide comprising anti-angiogenic alpha 1-24 peptide sequence derived from fibrinogen, useful for inhibiting angiogenesis sequence derived from fibrinogen, useful for inhibiting angiogenesis conditions such as cancer, neovascular glaucoma, psoriasis, rheumatoi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a polypeptide comprising an anti-angiogenic alpha 1-24 peptide sequence derived from fibrinogen. The peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Fig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (BIOA-) BIOACTA LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2002-339656/37.
DB; AAK99966.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 GPRVVERHOSA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GPRVVERHQSA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             78 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Lewis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; 2000GB-0027395.
; 2000US-252896P.
; 2001GB-0017737.
                                                                                                                                                                                                                                                                                                                             (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Score 57; Di
100.0%; Pred. No. 0.1
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Robinson
                                                                                                                                                                                                   noma; cancer; imaging agent; tumour growth inhibition;
                                                                                                                                                                                                                                                                                                                                                                                                  78 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ď
                                                                                                                                                                                     retinopathy;
                                                                                                                                                                     haemangioma; Kaposi's sarcoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 23;
0.0012;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                        plasmin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          rheumatoid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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14-MAY-2001; 2001WO-GB02079

22-NOV-2001

WO200188129-A1 Homo sapiens

Y)

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RABSULT 9
AAB5415
IID AAB54
XX AAB5
XX AAB5
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XX Huma
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XX Huma
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                                                                                                                                                                                                                                                                                                                                                        밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CC existing vascular bed, is a multistep process involving the degradation of components of the cellular matrix, leading to the migration, cc proliferation and differentiation of endothelial cells to form tubules of fibrin) is a dimeric molecule containing pairs of non-identical chains of ibrin) is a dimeric molecule containing pairs of non-identical chains of distrin) is a dimeric molecule containing pairs of non-identical chains of distrin is two outer D-domains and the central E-domain. Fibrinogen can be digested by either plasmin or thrombin, the fibrin E-fragment produced by proteolytic cleavage of fibrin stimulates angiogenesis in the correct in invasive breast carcinomas positively correlates with the degree of tumour vascularity. This invention has discovered proteolytic fragments of fibrinogen that inhibit angiogenesis. The nucleic acids and correcting cancer, and for treating a human or animal which would benefit from inhibition of angiogenesis. The proteins of the invention are useful in the manufacture of a medicament of or to monitor the effect of treatment to inhibit tumour growth). The coreins of the invention are also useful in modulating angiogenesis for the treatment of contains of the invention are also useful in modulating angiogenesis for the treatment of ophthalmological conditions (e.g. neovascular glaucoma, conditions (e.g. neovascular glaucoma, conditions (e.g. neovascular glaucoma, conditions (e.g. neovascular glaucoma, conditions of the invention are also useful in modulating angiogenesis for the treatment of ophthalmological conditions (e.g. neovascular glaucoma, conditions of the manufacture of a medicament conditions of the invention are also useful in modulating angiogenesis for the treatment of and the conditions (e.g. neovascular glaucoma, conditions of the invention are also useful in modulations (e.g. neovascular glaucoma, conditions of the conditions (e.g. neovascular glaucoma, conditions of the conditions of the condition of the conditions of the condition of the condition o
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local
Human; pancreas; pancreatic cancer; pancreatic cancer antigen; detection; diagnosis; identification; cytostatic; neuroprotective; nootropic; immunomodulatory; relaxant; contraceptive; ynaecological; antiinflammatory; cardiant; gene therapy; chromosome mapping;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel isolated nucleic acid molecule which encodes a fibrinogen polypeptide which is useful for treating cancer, diabetic retinc obesity, hepatitis, pneumonia, glomerulonephritis, asthma and
                                                                                                                                                           09-MAR-2001
                                                                                                                                                                                                                                             AAB54135 standard; Protein; 360
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         fibrinogen E-fragment polypeptides possessing anti-angiogenesis properties. Angiogenesis, the development of new blood vessels from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     E-fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      09-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13-MAY-2000;
14-JUN-2000;
                                                                                                               Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 6; Fig 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (UYSH-) UNIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   obesity);
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2002-062380/08
                                                                                                                                                                                                                                                                                                                                                        17
                                                                                                             pancreatic
                                                                                                                                                                                                                                                                                                                                                                                                 _
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                               GPRVVERHQSA 11
                                                                                                                                                                                                                                                                                                                                                        GPRVVERHQSA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (e.g. haemangioma, Kaposi's sarcoma,
/); inflammatory and infectious proces
and glomerulonephritis).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         alpha-chain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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2000GB-0014370.
2000GB-0027396.
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                                                                                                                                                        (first entry)
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                                                                                                             cancer antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       represents amino acids 1-78 in. The invention comprises I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         English
                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 57; DB 23;
Pred. No. 0.0012;
                                                                                                                                                                                                                                             B
                                                                                                           protein sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       comprises DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 processes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23;
                                                                                                                                                                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ovarian hyperstimulation
                                                                                                           SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           of the human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 78;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (e.g. hepatitis
                                                                                                             IJ
                                                                                                             NO:587
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           retinopathy,
                                                                                                                                                                                                                                                                                                                                                                                                                                          ..
..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              fibrinogen
                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              an
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cc gynaecological, cardiant and antiinflammatory activities, and can be used cin gene therapy. The polynucleotide and proteins can be used for preventing, treating, or ameliorating a medical condition or in assays cfor diagnosing a pathological condition or a susceptibility to one in a csubject. Binding partners to the proteins and the activity of the proteins can be identified. The pancreatic cancer antigens can be used to groteins can be identified. The pancreatic cancer antigens can be used to captible that can be screened for. The pancreatic can be used to captible that can be used to design nucleic candists and antagonists to the antigens can be screened for. The pancreatic cancer antigens can be used to design nucleic candispostic methods. The proteins can be used to generate antibodies can alysis, tissue identification and/or typing and a variety of forensic candispnostic methods. The proteins can be used to generate antibodies which are used to purify, detect and target the polypeptides, including continuous candinuous candinuous candisplant to proteins can be used to generate antibodies can be used to treat or prevent neural, immune system, muscular, reproductive, gastrointestinal, pulmonary, cardiovascular, renal or proteins can be used to treat or prevent neural, immune system, muscular, reproductive, disorders. AAC99220 to AAC99240 and AAE9467 represent
Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAC98773 to AAC99231 encode the human pancreatic cancer associated proteins, called pancreatic cancer antigens, given in AAB54008 to AAB54466. The human pancreatic cancer antigens have cytostatic, neuroprotective, nootropic, immunomodulatory, relaxant, contraceptive,
                                               sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 11; Page 1024-1026; 1379pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New nucleic acid that is a pancreatic cancer antigen for preventing, treating, or ameliorating a medical condition, particular pancreatic cancer, or for use in assays for diagnosing a pathological condition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2000-579444/54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rosen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         08-MAR-2000; 2000WO-US05989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200055320-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  linkage analysis; tissue identification; tissue typing; forensic; neural; immune system; muscular; reproductive; gastrointestinal; pulmonary; cardiovascular; renal; proliferative.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (HUMA-) HUMAN GENOME
                                               used in the
360
A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99US-0124270
                                               exemplification
                                               of the present
                                               invention
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                                                                                                                               Matches
                                                                                                                                     Query Match
Best Local (
Tissue binding;
tissue-binding o
                     Fibronectin
                                                                AAR60020
                                                                                                                                      Similarity
                                                                standard;
                                                                                                           GPRVVERHQSA
                                                                                                   GPRVVERHQSA
                                                                                                                               Conservative
                                    (first
tissue sealing; wound healing; vulnerary;
domain; TBD; crosslinking domain; fibronectin;
                                                              Protein; 643
                                                                                                                                     100
                                                                                                   67
                                                                                                                                     0.0%;
                                                                                                                               0
                                                                                                                                     Score 57;
Pred. No.
                                                                                                                               Mismatches
                                                                                                                                     DB 21;
                                                                                                                               0
                                                                                                                                           Length
                                                                                                                               Indels
                                                                                                                                             360;
                                                                                                                               0
                                                                                                                               Gaps
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domain;

Prunkard

by introducing l and collecting collecting

DNA

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Human fibrinogen A-alpha chain; transgenic animal milk; treatment; sheep beta-lactoglobulin signal peptide fusion protein; surgical adhesive.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hybrid proteins have a tissue-binding domain (TBD) from 1 protein linked to a crosslinking domain from another protein. The TBD given in AAR6019; the heparin-binding domain (aa 1812-2171 of AAR60021) of fibronectin; the collagen-binding domain (aa 282-608 of AAR60021) of fibronectin; or the cell-binding domain (aa 387-1903 or 1532-1631 of AAR60020) of fibronectin. DNA encoding a fibronectin-fibrinogen hybrid is given in AAQ70007, and sequences for fibronectin and fibrinogen-alpha in AAQ70008 and AAQ70009, respectively.
                                                                                                                                                                                                                                                                                                                                     AAR82244 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 57-60; 87pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New hybrid proteins for use in tissue sealing and wound healing comprising a tissue-binding domain from a protein covalently linked to a crosslinking domain of another protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1994-249231/30.
N-PSDB; AAQ70008.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (ZYMO ) ZYMOGENETICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-DEC-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-DEC-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21-JUL-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO9416085-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     fibrinogen-alpha; heparin-binding domain; collagen-binding domain; cell-binding domain; hybrid protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                     fibrinogen A-alpha chain protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       36 ĠPŔVVĖŔĦQŚŔ 46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 GPRVVERHOSA 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               643 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   92US-0998271.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                93WO-US12687.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note= "acts as tissue-binding protein"
                                                                                                                                                                                                                                                                                                                                   Protein; 644 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /label=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      label=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; ;
100.0%; ;
itive 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      = cell-binding_domain
"acts as tissue-binding domain of hybrid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cell-binding_domain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 57; DB 15;
Pred. No. 0.012;
Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 643;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        domain of hybrid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
RESULT 12
AAY82891
XX AAY82
XX AAY82
XX 19-JU
XX 19-JU
XX 19-JU
XX Fibri:
KW Fibri:
KW uroki
KW myoca
KW myoca
KW MO200
XX Homo
PN WO200
XX 12-AU
XX 12-AU
XX 12-AU
XX (NYBL
X
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Best Local S
Matches 11
                                                          (NYBL-) NEW YORK BLOOD CENT
                                                                                                               12-AUG-1998;
                                                                                                                                                            12-AUG-1999;
                                                                                                                                                                                                          24-FEB-2000
                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                      Fibrinogen; blood; blood clotting; plasminogen activator; urokinase; streptokinase; alpha subunit; beta subunit; gamma subunit; myocardial infarction; reperfusion; atherogenesis; alphaEC; alphaECX; human; trauma; wound repair; angiogenesis; cancer.
                                                                                                                                                                                                                                                          WO200009562-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    AlphaE subunit of human fibrinogen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAY82891;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAY82891 standard; Protein; 847 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human fibrinogen A-alpha chain may be produced recombinantly, preferably by sheep beta-lactoglobulin signal peptide fusion protein gene expression in non-human transgenic animal milk, fibrinogen B-beta chain (see AAR82243) and the gamma chain (see useful in human and veterinary medicine, e.g. in the formulation of surject alachesives, which also consist of Factor-XIII, and as a coating surface for polymeric articles, such as synthetic vascular grafts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 42-44; 99pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Production of fibrinogen in transgenic mammals - 1 segments into the germ line of a non-human mammal milk from female progeny.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1995-320582/41.
N-PSDB; AAT03853.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Dalrymple MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    03-MAR-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-MAR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  08-SEP-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (PHAR-) PHARM PROTEINS LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO9523868-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 GPRVVERHOSA 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GPRVVERHOSA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 644 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                          98US-0096210
                                                                                                                                                     99WO-US18412
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               9408-0206176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Foster DC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             95WO-US02648
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 57; DB 16;
Pred. No. 0.012;
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Mismatches

0,

0,

Gaps

0,

Length 644;

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Domain

Domain Key

RESULT 11
AAR82244
ID AAR82
XX AAR82
XX AAR82
XX DT 26-OC
XX Uman
XX Human
XX Human
XX Sheep
XX Sheep
XX Sheep
XX Sheep
XX Sheep
XX Hombo

Human 26-OCT-1996 AAR82244;

Homo sapiens

Grieninger G,

Applegate D,

Stoike-steben L;

В Ş

Query Match Best Local S Matches 11

11;

Sequence

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RESULT 13
AAW11932
conditions associated with fibrinogen metabolism e.g. trauma
conditions associated with fibrinogen metabolism e.g. trauma
crequiring wound repair, angiogenesis and cancer. They are useful
conditions associated with fibrinogen alphaECX domain in vivo or in
control the presence of alphaECX cleavage fragments indicates
control the presence of alphaECX cleavage fragments indicates
control to the presence of alphaECX cleavage fragments indicates
control to the presence of alphaECX cleavage fragments indicates
control to the subclass of human
control to the sub
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      밁
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ribrinogen has a complex structure comprising two copies of alpha, beta and gamma subunits in a heavily disulfide bonded hexamer. Until recently it was thought that the alpha subunits lacked a globular C-terminal domain which was present on the beta and gamma subunits. However, an additional exon (exon VI) downstream of the alpha subunit gene and additional splicing leads to the use of this extra sequence to elongate the alpha chain by 35% (236 amino acids), providing the alpha subunit with a globular C-terminal domain. Isolated fragments
                                                                                                                                                                                     01-NOV-1994;
                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             disseminated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Monoclonal; antibody; human; soluble; fibrin; fibrinogen; urea-treated; des-AABB; B-beta-chain; immunoassay; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         of human fibrinogen termed 'alphaECX cleavage fragments' derived from this globular C-terminal domain can be administered therapeutically to treat mammals (especially humans) suffering from the transfer of the treatment of the 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New fragments of human fibrinogen, useful for treating conditions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-205983/18.
N-PSDB; AAZ93039.
                                                                                                          02-NOV-1993;
                                                                                                                                                                                                                                                                11-MAY-1995
                                                                                                                                                                                                                                                                                                                                           WO9512617-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAW11932 standard; peptide; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      associated with fibrinogen metabolism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Fibrinogen B-beta-chain residues 15-24.
                                   (IATR ) IATRON LAB INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         intravascular coagulation
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0.017;
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RESULT 14
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Best Local S
Matches 10
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                                                                                                     New purified luciferase polypeptide, useful for identifying protease activity modulators, comprises recognition site cleavable by protease where cleavage of polypeptide results in decreased luciferase activity.
                                                                                                                                                                                                                                                                                                                          Luciferase; cell death;
                                                                                                                                                                      Leng
                                                                                                                                                                                                                                                                                                                                                           Thrombin
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                                                                                                                                                 WPI; 2002-195809/25.
                                                                                                                                                                                                                 19-JUL-2000; 2000US-0619047
                                                                                                                                                                                                                                     17-JUL-2001; 2001WO-US22478
                                                                                                                                                                                                                                                            24-JAN-2002.
                                                                                                                                                                                                                                                                                WO200206458-A2
                                                                                                                                                                                                                                                                                                      Unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                      AAE19125;
                                                                                                                                                                                                                                                                                                                                                                                                                          AAE19125 standard; peptide; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         disseminated intravascular coagulation.
                                                                                                                                                                                            (CHEM-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10;
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                                                                                                                                                                                                                                                                                                                           enzyme; p
thrombin.
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The invention relates to a purified polypeptide having luciferase activity and a recognition site specifically cleavable by a protease, where cleavage results in a decrease in luciferase activity. The polypeptide comprises a localisation sequence which is linked to the luciferase polypeptide by the cleavable recognition sequence. The polypeptide is useful for identifving a nearest form.

Claim 63;

Page

24; 56pp;

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В
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a polypeptide comprising an anti-angiogenic alpha 1-24 peptide sequence derived from fibrinogen. The peptide comprising ADSGEGDFLAEGGGVRGPRVVERH (alpha 1-24 peptide) is useful for manufacturing a medicament for treating cancer. The peptide, its encoding DNA, a pharmaceutical composition and vector of the invention are useful for treating an animal which would benefit from inhibition of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity .100.0%;
Matches 8; Conservative (
                                                                                                                               Claim 2;
                                                                                                                                                Novel polypeptide comprising anti-angiogenic alpha 1-24 peptide sequence derived from fibrinogen, useful for inhibiting angiogenesis in conditions such as cancer, neovascular glaucoma, psoriasis, rheumatoid
                                                                                                                                                                                                                                                               Staton C,
                                                                                                                                                                                                                                                                                                                 01-SEP-2000; 2000GB-0021475.
09-NOV-2000; 2000GB-0027395.
27-NOV-2000; 2000US-252896P.
23-JUL-2001; 2001GB-0017737.
                                                                                                                                                                                                                                                                                                                                                                                           03-SEP-2001; 2001WO-GB03926
                                                                                                                                                                                                                                                                                         (BIOA-) BIOACTA LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                               07-MAR-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200218440-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cytostatic; antitumour; ophthalmological; antiarthritic; antirheumatic; antipsoriatic; antidiabetic; anti-anglogenic; alphal-24 peptide; tumour fibrinogen; cancer; angiogenesis; chemotherapeutic agent; psoriasis; neovascular glaucoma; rheumatoid arthritis; diabetic retinopathy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Anti-angiogenic alpha 1-24 peptide #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAO21112 standard; Peptide; 24 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                modulator, an inhibitor of apoptosis and for detecting luciferase activity in a sample. The polypeptide is used for characterising and identifying cellular processes associated with metabolism, cell growth protease measurement methods are useful for characterising, identifying cellular biochemical pathways as well as identifying diagnostic and siecemical pathways as well as identifying diagnostic and biochemical pathways as well as identifying diagnostic and study agents or conditions that cleave the recognition site and for activity of protease inside a cell that expresses a recombinant invention. The present sequence is thrombin peptide used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                               2002-339656/37.
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                                                                                                                                                                                                                                                           Lewis C,
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Best Local
                                                                                                                                      angiogenesis, where the therapeutic agents are administered to the animal and the effects of the agent on the inhibition of angiogenesis is monitored. Polypeptides useful for treatment are conjugated, associated or crosslinked to a chemotherapeutic agent. The method is preferably its encoding DNA, a pharmaceutical composition and vector of the invention are useful for inhibiting angiogenesis in tumours. The peptide, invention are useful for inhibiting angiogenesis in conditions of retinopathy. The DNA of the alpha 1-24 peptide can be used in gene therapy. This sequence represents an anti-angiogenic alpha 1-24 peptide of the invention.
                                                                                                                             Sequence
17 GPRVVERH
                         1 GPRVVERH
                                                              8;
                                                                             Similarity
                                                                                                                          24 AA;
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Search completed: June 27, Job time : 34.55 secs 2003, 17:16:47

Copyright

GenCore version (c) 1993 - 2003

5.1.6 Compugen Ltd.

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Minimum
Maximum
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57
1 GPRVVERHQSA 11
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| G972_6/ptodata/1/pubpaa/US08_NEW
| G972_6/ptodata/1/pubpaa/US06_NEW
| G972_6/ptodata/1/pubpaa/US06_NEW
       GPRVVERHQSA 11
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cgn2_6/ptodata/1/pubpaa/US06/
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US-09-931-009A-1
US-09-919-039-121
US-10-112-527-4
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US-10-062-831-106
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NEW PUB.pep:*
6 NEW PUB.pep:*
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Sequence 11, Appl
Sequence 10, Appli
Sequence 7, Appli
Sequence 190, Appli
Sequence 2, Appli
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Sequence 216, Appl
Sequence 106, Appl
Sequence 106, Appl
Sequence 79, Appl
Sequence 11695, A
Sequence 11695, A
Sequence 486, Appl
Sequence 486, Appl
Sequence 678, Appl
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Sequence 121,
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equence 1, Appli
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GENERAL INFORMA
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ALIGNMENTS

RESULT 1 US-09-925-297-587 ; Sequence 587, A

Sequence 587, Application US/09925297 Patent No. US20020081659A1 GENERAL INFORMATION:

Antibodies

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; OTHER INFORMATION: US-09-925-297-587
                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO 587
                                                                                                                                                                                                                                                                                              LENGTH: 360
TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: SITE
LOCATION: (15)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT FILING DATE: 2001-08-10.
PRIOR APPLICATION NUMBER: PCT/US00/05989
PRIOR FILING DATE: 2000-03-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins and
FILE REFERENCE: PA105
CURRENT APPLICATION NUMBER: US/09/925,297
CURRENT FILING DATE: 2001-08-10
                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 928
SOFTWARE: PatentIn Ver. 2.0
                                                                                  LOCATION: (326)
OTHER INFORMATION: :
NAME/KEY: SITE
LOCATION: (327)
                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: 60/124,270 PRIOR FILING DATE: 1999-03-12
                                  OTHER INFORMATION:
NAME/KEY: SITE
LOCATION: (339)
                                                                                                                                                                                                                          NAME/KEY: SITE
LOCATION: (315)
OTHER INFORMATION:
                                                                                                                                                       NAME/KEY: SITE
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                Xaa equals any of the naturally occurring L-amino acids
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Query Match

100.0%; Score 57;

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RESULT 4
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US-09-931-009A-J
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                 Sequence 4, Application US/10112527
Patent No. US20020168722A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/919,039
CURRENT FILING DATE: 2002-09-09
PRIOR APPLICATION NUMBER: 60/222,113
PRIOR FILING DATE: 2000-07-28
NUMBER OF SEQ ID NOS: 401
SOFTWARE: PERL PROGRAM
SEQ ID NO 121
LENGTH: 644
                                                                                                                                                                                                                      Query Match
Best Local :
                                                                                                                                                                                                       Matches
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Best Local Similarity
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                                                                                                                                                                                                                                                                     -09-919-039-121
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APPLICANT: Grieninger, Gerd
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Kaser, Matthew R.
TITLE OF INVENTION: GENES EXPRESSED IN TREATED HUMAN C3A LIVER CELL CULTURES
FILE REFERENCE: PA-0035 US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EQ ID NO 1
LENGTH: 620
TYPE: PRT
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CURRENT APPLICATION NUMBER: US/09/931,009A
CURRENT FILING DATE: 2002-06-28
                                                                                                                                                                                                                                                                              NAME/KEY: misc_feature
OTHER INFORMATION: Incyte ID No. US20030108871A1 280276CD1
                                                                                                                                                                                                                                                                                                                                  ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 2
                                                                                                                                                                                                                                                                                                                    FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Homo sapiens
                                                                                                                                                                                                                   Match 100.0%;
Local Similarity 100.0%;
                                                                                                                                   36 GPRVVERHOSA 46
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Pred. No. 0.017;
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                                                                                                                                                                                                                             DB 9; Length 644;
                                                                                                                                                                                                  0; Indels
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Query Match
Best Local Similarity
"hes 8; Conserva
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US-09-953-657-5
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Best Local (
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CURRENT FILING DATE: 2002-03-29
PRIOR APPLICATION NUMBER: US 09/373,157
PRIOR FILING DATE: 1999-08-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn version 3.1
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                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: 08/595,718
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 17-Sep-2001
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                   SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette,
                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Baumbach, George A.
Hammond, David J.
TITLE OF INVENTION: Fibrinogen Binding Peptides
NUMBER OF SEQUENCES: 8
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                                                                                                                                                                                                      TELEPHONE: (510)705-7910
TELEFAX: (510)705-7904
                                                                                                                                                                                                                                                  NAME: Giblin, James A.
REGISTRATION NUMBER: 25772
REFERENCE/DOCKET NUMBER: MSB-7233
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 800 Dwight Way
P. O. Box 1986
                                                                                                                                                                          VOLUME:
                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/09/953,657
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                                                                                                                                    439-444
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            78.9%; Score 45; DB 10;
88.9%; Pred. No. 3.8e+05;
tive 1; Mismatches 0;
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Pred. No.
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                                              Length 9;
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US-10-057-505-11

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                                      ; OTHER INFORMATION: US-09-765-614B-10
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                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 31
SOFTWARE: PatentIn Ver.
SEQ ID NO 10
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APPLICANT: Nycomed Imaging AS
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Best Local
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Patent No. US20020102215A1
                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: REF/Klaveness/054
CURRENT APPLICATION NUMBER: US/09/765,614B
CURRENT FILING DATE: 2001-07-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: US 09/396,003
PRIOR FILING DATE: 1999-09-13
NUMBER OF SEQ ID NOS: 31
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/10/057,505
CURRENT FILING DATE: 2002-01-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: CUBITT, Andrew
TITLE OF INVENTION: TANDEM FLUORESCENT PROTEIN CONSTRUCTS
FILE REFERENCE: REGEN1260-3
                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: Improvements in or relating to TITLE OF INVENTION: diagnostic/therapeutic TITLE OF INVENTION: agents
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                                                                                                                                                                                                                           TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                            NAME/KEY: MOD_RES
LOCATION: (1)
OTHER INFORMATION: Biotinylated-Gly
                                                                                                                                                                    OTHER INFORMATION: Description of Artificial OTHER INFORMATION: Sequence:Biotinylated OTHER INFORMATION: fibrin-anti-polymerant peptide
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                                                                                              NAME/KEY: MOD_RES
                                                                            OCATION:
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TSIEN, Roger
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                                                          AMIDATION
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  71.9%;
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Pred. No.
Score 41;
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  DB 10;
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Length 10;
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US-10-205-823-190
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Matches
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Patent No. US20020102217A1
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APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                    Publication No.
                                                                                                                     APPLICANT: Anderson, Dustin
TITLE OF INVENTION: MOVEL GENES, COMPOSIT
TITLE OF INVENTION: METHODS FOR IDENTIFI
TITLE OF INVENTION: THERAPY OF PROSTATE
FILE REFERENCE: MRI-044
                                                                                                                                                                                                  APPLICANT:
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CURRENT APPLICATION NUMBER: US/10/205,823
CURRENT FILING DATE: 2002-07-25
PRIOR APPLICATION NUMBER: 60/307,982
PRIOR FILING DATE: 2001-07-25
PRIOR APPLICATION NUMBER: 60/314,356
PRIOR FILING DATE: 2001-08-22
PRIOR APPLICATION NUMBER: 60/325,020
PRIOR APPLICATION NUMBER: 60/325,020
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/925,715
CURRENT FILING DATE: 2001-08-10
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                                                                                                                                                                                                                                       APPLICANT:
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LOCATION: (1)
OTHER INFORMATION: Biotin-Gly
NAME/KEY: MOD RES
LOCATION: (10)
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: AMIDATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: Description of Artificial Sequence:Biotinylated OTHER INFORMATION: fibrin-antipolymerant peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ENGTH: 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GPRVVERHQS 10
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                                                                                                                                                                                                                                                  Hoersch, Sebastian
Kamatkar, Shubhangi
Wonsey, Angela M.
                                                                                                                                                                                                                                                                                                        Gannavarapu, Manjula
Gorbatcheva, Bella
                                                                                                                                                                                                                                                                                                                                              Endege, Wilson O.
                                                                                                                                                                                                                                                                                                                                                                                                                                   Application US/10205823
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                                                                                                                                                                                                                  Xumei
                                                                                                                                                                                                                                       Karen
                                                                                                                                                                                                                                                                                                                                                                John E
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 41;
Pred. No.
                                                                                                                                                                                 COMPOSITIONS,
                                                                                                                                                                IDENTIFICATION,
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APPLICANT: LUPRATE, Fabrice
APPLICANT: Lesage, Florian
APPLICANT: Lesage, Florian
APPLICANT: Lesage, Florian
APPLICANT: Lesage, Florian
APPLICANT: Lazdunski, Michel
ITITE OF INVENTION: FAMILY OF MAMMALIAN POTASSIUM CHANNELS, THEIR CLONING
FILE REFERENCE: 1201-CIP-DIV-00
CURRENT APPLICATION NUMBER: US/09/939,484
FRICA REPLICATION NUMBER: US/09/939,484
PRIOR APPLICATION NUMBER: 09/144,914
PRIOR APPLICATION NUMBER: 09/144,914
PRIOR FILING DATE: 1998-09-01
PRIOR APPLICATION NUMBER: 08/749,816
PRIOR FILING DATE: 1998-09-01
PRIOR FILING DATE: 1998-08-08
PRIOR FILING DATE: 1998-08-08
PRIOR FILING DATE: 1998-08-08
PRIOR FILING DATE: 1998-08-08
PRIOR FILING DATE: 1998-02-08
NUMBER: OF SEQ ID NOS: 24
SCOTTWARDE: DEFAULT TOWN
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Best Local S
APPLICANT: Duprat, Fabrice
APPLICANT: Duprat, Fabrice
APPLICANT: Lesage, Florian
APPLICANT: Eink, Michel
APPLICANT: Lazdunski, Michel
TITLE OF UNVENTION: FAMILY OF MAMMALIAN POTASSIUM CHANNELS, THEIR CLONING
TITLE OF INVENTION: AND THEIR USB, ESPECIALLY FOR THE SCREENING OF DRUGS
                                                                                                                               Sequence 2, Application US/09939483 Patent No. US20020094558A1 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO 2
LENGTH: 336
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Patent No. US20020032322A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       -09-939-484-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: FastSEQ for Windows Version SEQ ID NO 190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: TWIK-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 60/341,746
PRIOR FILING DATE: 2001-12-12
PRIOR APPLICATION NUMBER: 60/362,158
PRIOR: FILING DATE: 2002-03-05
NUMBER OF SEQ ID NOS: 455
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77.8%;
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Pred. No.
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Pred. No. 4
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Query Match
Best Local Similarity
Wortches 6; Conserve
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; ORGANISM: Escherichia coli
US-10-025-222A-22
Sequence 106, Application US/10062831
Publication No. US20030105297A1
GENERAL INFORMATION:
APPLICANT: Steven M. Ruben, et al.
TITLE OF INVENTION: 32 Human Secreted Proteins
FILE REFERENCE: PZ006P1
CURRENT APPLICATION NUMBER: US/10/062,831
CURRENT FILING DATE: 2002-02-05
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US-10-025-222A-22
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LENGTH: 581
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Best Local S
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PRIOR FILLING DATE: 1998-09-01
PRIOR APPLICATION NUMBER: 08/749,816
PRIOR APPLICATION NUMBER: 60/095,234
PRIOR APPLICATION NUMBER: 60/095,234
PRIOR FILLING DATE: 1998-08-04
PRIOR APPLICATION NUMBER: FR 96/01565
PRIOR APPLICATION NUMBER: FR 96/01565
PRIOR FILLING DATE: 1996-02-08
NUMBER OF SEQ ID NOS: 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: 60/256,349
PRIOR FILING DATE: 2000-12-19
NUMBER OF SEQ ID NOS: 41
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: DELEGTIER, JERRY
APPLICANT: DELEGTIER, JERRY
APPLICANT: DELEGTIER, MICHAEL
TITLE OF INVENTION: COMPOSITIONS AND METHODS INVOLVING AN ESSENTIAL
TITLE OF INVENTION: STAPHYLOCOCCUS AUREUS GENE AND ITS ENCODED PROTEIN
FILE OF INVENTION: STAPHYLOCOCCUS AUREUS GENE AND ITS ENCODED PROTEIN
FILE REFERENCE: 073406-0771
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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SEQ ID NO 2
LENGTH: 336
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/10/025,222A CURRENT FILING DATE: 2001-12-19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
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O. US20030003444A1
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Pred. No. 1
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Pred. No.
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-062-831-106
                                                                                                                                                                                                       CURRENT FILING DATE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR FILING DATE: 2000-02-04
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR APPLICATION NUMBER: US 20263.6
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-10-04
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Best Local S
Matches 5
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SEQ ID NO 106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 46461, Application US/09864761 Patent No. US20020048763A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Penn, Sharron G. APPLICANT: Rank, David R.
                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT FILING DATE: 2001-05-23
                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY FILE REFERENCE: Acomica-X-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OR APPLICATION NUMBER: 60/056,250
OR FILLING DATE: August 29, 1997
OR APPLICATION NUMBER: 60/056,296
OR FILLING DATE: August 29, 1997
OR APPLICATION NUMBER: 60/056,293
OR FILLING DATE: August 29, 1997
                                   APPLICATION NUMBER: PCT/US01/00664
FILING DATE: 2001-01-30
                                                                                                                                            APPLICATION NUMBER: US 60/236,359
FILING DATE: 2000-09-27
APPLICATION NUMBER: PCT/US01/00666
                                                                                   APPLICATION NUMBER: PCT/US01/00667 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00669 FILING DATE: 2001-01-30
                                                                                                                            FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: May 30, 1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: MAY 30, 1997
APPLICATION NUMBER: 60/048,356
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APPLICATION NUMBER: 60/048,101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: May 30, 1997
APPLICATION NUMBER: 60/050,935
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APPLICATION NUMBER: 60/048,190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 60/048,093
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: May 28, 1998
APPLICATION NUMBER: 60/044,039
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
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, David K.
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Pred. No. 17;
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PRIOR APPLICATION UNMBER: PCT/US98/10868
PRIOR APPLICATION UNMBER: PCT/US98/10868
PRIOR APPLICATION UNMBER: 60/044,039
PRIOR FILING DATE: May 28, 1998
PRIOR APPLICATION UNMBER: 60/048,039
PRIOR APPLICATION UNMBER: 60/048,093
PRIOR APPLICATION UNMBER: 60/048,190
PRIOR APPLICATION UNMBER: 60/048,190
PRIOR APPLICATION UNMBER: 60/050,935
PRIOR FILING DATE: May 30, 1997
PRIOR FILING DATE: May 30, 1997
PRIOR APPLICATION UNMBER: 60/048,101
PRIOR APPLICATION UNMBER: 60/048,101
PRIOR APPLICATION UNMBER: 60/048,101
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US-10-062-831-79
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Best Local S
Matches 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 79, Appublication No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/10/062,831
CURRENT FILING DATE: 2002-02-05
PRIOR APPLICATION NUMBER: 09/690,454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: 32 FILE REFERENCE: PZ006P1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Steven M. Ruben,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE:
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OTHER INFORMATION: EXPRESSED IN ADULT
OTHER INFORMATION: EXPRESSED IN FETAL
OTHER INFORMATION: EST_HUMAN HIT: AV
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TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
                                        OR FILING DATE: May 30, 1997

OR APPLICATION NUMBER: 60/046,356

OR FILING DATE: May 30, 1997

OR APPLICATION NUMBER: 60/056,250

OR FILING DATE: August 29, 1997

OR APPLICATION NUMBER: 60/056,250

OR FILING DATE: August 29, 1997

OR FILING DATE: August 29, 1997
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APPLICATION
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: PCT/US01/00668 FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00661
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                                 N NUMBER: 60/056,296
N NUMBER: 60/056,296
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SCOTUMARE: PatentIn Ver. 2.0
; SEQ ID NO 79
; LENGTH: 208
; TYPE: PAT
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: SITE
; LOCATION: (148)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
; PEATURE:
; NAME/KEY: SITE
; LOCATION: (186)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
; PEATURE:
; NAME/KEY: SITE
; LOCATION: (208)
; OTHER INFORMATION: Xaa equals stop translation
US-10-062-831-79

Query Match
Best Local Similarity 55.6%; Pred. No. 91;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

Dy 2 PRVVERHQS 10
ph 62 PRILEAHQN 70

Search completed: June 27, 2003, 17:30:21

Job time: 35.1 secs
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Minimum
Maximum
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Perfect score:
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Maximum Match 100%
Listing first 45 summaries
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     Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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/cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
/cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
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 US-08-595-718A-5
US-08-950-054A-10
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US-08-959-206A-7
US-08-959-206A-7
US-08-959-206A-7
US-08-959-200-40
US-08-195-006-40
US-08-195-006-40
US-08-195-006-40
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US-08-195-006-40
US-08-958-408B-25
US-09-257-581-5
US-09-257-581-7
52604323-3
US-09-117-853-2
US-08-764-100-27
US-08-764-100-27
US-08-96-338-20
US-08-96-338-20
US-08-96-338-20
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US-08-551-356-4

PCT-US93-12687-4

US-08-206-176-2

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US-08-448-547-2
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Sequence 7, Appl
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Sequence 11, Appl
Sequence 10, Appl
Sequence 40, Appl
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Sequence 25, Appl
Sequence 5, Appl
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Sequence 27, Appl
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54.4	54.4	54.4		54.4			54.4	56.1	56.1	56.1		56.1	56.1	56.1	56.1	56.1	56.1
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US-08-507-016-4	US-08-507-016-3	PCT-US95-13794-52	US-08-469-141A-52	PCT-US95-13794-24	US-08-469-141A-24	US-08-860-808E-1	US-08-434-099A-37	US-07-945-283-2	US-08-808-599A-5	US-08-728-626-5	US-08-738-975-5	US-08-751-965-5	US-08-439-818A-5	US-08-317-522A-5	US-09-578-178-2	US-09-173-151A-28	US-09-110-618-2
Sequence 4, Appli	Sequence 3, Appli	Sequence 52, Appl	Sequence 52, Appl	Sequence 24, Appl	Sequence 24, Appl	Sequence 1, Appli	Sequence 37, Appl		•	Sequence 5, Appli	-	-	•	•	Sequence 2, Appli	Sequence 28, Appl	Sequence 2, Appli

ALIGNMENTS

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RESULT 2
US-08-551-356-4
; Sequence 4, Application
; Patent No. 5830700
; PATENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Memon-Tudolph, Sunitha
APPLICANT: Chosh-Dastidar, Pradig
FITTLE OF INVENTION: Polypeptides Having a Single
FITTLE OF INVENTION: N-Terminal Water-Soluble Pol
TITLE OF INVENTION: Compositions and Kits
FILE REFERENCE: SEQUENCE LISTING ORT 843
CURRENT APPLICATION NUMBER: US/08/905,310
CURRENT FILING DATE: 1997-08-01
NUMBER OF SEQ ID NOS: 12
SOPTWARE: Patentin Ver. 2.0
SEQ ID NO 12
LENGTH: 13
TYPE: PRT
ORGANISM: Unknown
FEATURE: THE ORGANISM: Unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; OTHER INFORMATION: Description of Unknown Organism:peptide fragment US-08-905-310-12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 100.0%;
Best Local Similarity 100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Wei, Ziping
ZIP: 98105
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                             APPLICANT: Irani, Meher TITLE OF INVENTION: HYBI
                                                                                                                                                       NUMBER OF SEQUENCES: 1
                                                           STATE: WA
COUNTRY: USA
                                                                                               STREET: 4225 F
                                                                                                                                       ADDRESSEE:
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                                                                                                                                                                                                                                                                       Application US/08551356
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                                                                                                                   Roosevelt Way, N.E
                                                                                                                                                                                               HYBRID CROSS-LINKING PROTEINS
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Pred. No. 5.5e-05;
Pred. No. 5.5e-05;
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                                INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                  ZIP: 98105
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
SEQUENCE CHARACTERISTICS:
LENGTH: 643 amino acids
                                                                                                                                                                                                                                                            COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PATENTIN Release #1.0, CURRENT APPLICATION DATA:
                                                              TELECOMMUNICATION INFORMATION:
TELEPHONE: 206-547-8080 ext 322
                                                                                                                                              FILING DATE: 31-DEC-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: protein
                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: 92-26
TELECOMMUNICATION INFORMATION:
TELEPHONE: 206-547-8080 ext 322
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PRIOR APPLICATION DATA:
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                                                                                              NAME: Parker, Gary E
REGISTRATION NUMBER: 31
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity hes 11; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                      CITY: Seattle
STATE: WA
                                                  TELEFAX:
                                                                                                                                                                                                                CLASSIFICATION:
                                                                                                                                                                                                                                 FILING DATE:
                                                                                                                                  NAME:
                                                                                                                                                                                                                                              APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Holly, Julie A REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/07/998,271 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     36 GPRVVERHOSA 46
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4225 Roosevelt Way, N.E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               643 amino acids
                                                  206-548-2329
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HYBRID CROSS-LINKING PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%;
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                                                                                              92-26PC
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Pred. No. 0.0033;
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                                                                                                                                                                                                                                                                           Version
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Best Local S
Matches '11
                                              GENERAL
                                                          Sequence 4, Application US/09373157 Patent No. 6416963
                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                     Query Match
Best Local :
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APPLICANT: Grieninger, Gerd
APPLICANT: Applegate, Dianne
APPLICANT: Stoike-Steben, Lara
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Patent No.
                                                                                                                                                                                                                                                                                                                                  TELEFAX: 206-548-2329
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMAPPLICANT:
                                                                                                                                                                                                                                                                                                                                                                     NAME: Parker, Gary E
REGISTRATION NUMBER: 31-648
REFERENCE/DOCKET NUMBER: 93-15
TELECOMMUNICATION INFORMATION:
TELEPHONE: 206-547-8080 ext 322
                                                                                                                                                                                                                                                                         TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ALUNCA STREET: 42.
STREET: 42.
CITY: Seattle
TITE: WA
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MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                                                                                TELEPHONE: 200-548-2329
                                                                                                                                                                                                                                                                                                       TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                   LENGTH:
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                                             INFORMATION:
                                                                                                                                              36 GPRVVERHOSA 46
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                                                                                                                                                                                                                                                                                                                     644 amino acids
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4225 Roosevelt Way, N.E.
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                                                                                                                                                                                                      100.0%; ilarity 100.0%; Conservative 0;
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Prunkard, Donna E
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Pred. No. 0.0033;
; Mismatches 0
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                                                                                                                                                                                                                               DB 1; Length 644;
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                                   ; MOLECULE TYPE: peptide US-08-448-547-2
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; ORGANISM: Homo sapiens
US-09-373-157-4
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CURRENT FILING DATE: 1999-08-12
NUMBER OF SEQ ID NOS: 7
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 4
   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches 11; Conservative
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                              TELEFAX: (202)293-7860
NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION DATA:
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                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: Q-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)293-7060
                                                                                                                                                                                                                                                                    ATTORNEY AGENT INFORMATION:
NAME: NAKAMURA, Dean H.
                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: PCT/JP94/01844
FILING DATE: 01-NOV-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 5-297325
                                                                                                                                               SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: UFILING DATE: 30-JUN-1
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
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APPLICANT: Inuzuka, Kimiko
APPLICANT: Ito, Yumiko
TITLE OF INVENTION: ANTI-HUMAN SOLUBLE FIBRIN ANTIBODY,
TITLE OF INVENTION: HYBRIDONA, AND IMMUNOASSAYING METHOD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Washington STATE: D.C.
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                                                                         TOPOLOGY:
                                                                                      STRANDEDNESS:
                                                                                                                                                                                                                                                   REGISTRATION NUMBER: 33,981
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                                                                                                                            ENGTH:
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                                                                                                           amino acid
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                                                                                                                          10 amino acids
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Kohno,
                                                                         linear
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 93.0%;
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 Score 53;
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DB 2;
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Length 10;
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; DATE: 28-JUN-1990
US-08-595-718A-5
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     Matches
                                    Query Match
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INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                    PUBLICATION INFORMATION:
AUTHORS: Kuyas, C.
AUTHORS: Haeberli, A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: WordPerfect 6.1 CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE:
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STRANDEDNESS: SING
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y Match 78.9%; Score 45; DB 1; Length 9;
Local Similarity 88.9%; Pred. No. 1.9e+05;
hes 8; Conservative 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Giblin, James A.
REGISTRATION NUMBER: 2577
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE:
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                                                                                                                                                                                                                                                                                                         DESCRIPTION:
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                                                                                                                       SSUE:
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                                                                                                                                                   B: Walder, P. W.
B: Straub, P. W.
B: Straub, P. W.
B: Isolation of Human Fibrinogen and its
Derivatives by Affinity Chromatography on Gly-Pro-Arg-Pro-Lys-Fractogel
B: Thromb. Haemost.
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NVENTION: Fibrinogen Binding Peptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              E: Bayer Corporation
800 Dwight Way
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 IBM
                                                                                                                                                                                                                                                                                                                                                                                                                                                           (510)705-7910
                                                                                                                                                                                                                                                                                                         peptide
                                                                                                                                                                                                                                                                                                                                                       single strand
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mismatches 0;
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                                NUMBER: PAI
SOFTWARE: PAI
SEQ ID NO 10
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                                                                                                                                                                                                                                                                                                                                                                                                                ; MOLECULE TYPE: peptide
US-08-792-553-11
                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                         Sequence 10,
                                                                                 TITLE OF INVENTION: Improvements in or rel
TITLE OF INVENTION: diagnostic/therapeutic
FILE REFERENCE: REF/Klaveness/054
CURRENT APPLICATION NUMBER: US/08/960,054A
CURRENT FILING DATE: 1997-10-29
                                                                                                                                                                                                            Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPAX: 619-678-5099
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 17 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
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                                                                        NUMBER OF SEQ ID NOS:
                                                                                                                                                                               APPLICANT:
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Lisa A. Haile, Ph.D..
REGISTRATION NUMBER: 38,347
REFERENCE/DOCKET NUMBER: 077
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 31-JAN-1997
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PATENTIN Rel
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: 31-JAN-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: La Jolla
STATE: California
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10. 5981200
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                                                             PatentIn Ver.
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Heim, Roger
VENTION: Tandem Fluorescent Protein Constructs
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                                                                                                                                          diagnostic/therapeutic
                                                                                                                                                      Improvements in or relating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Release #1.0, Version #1.25
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APPLICANT: Nycomed Imaging AS
ITITLE OF INVENTION: Improvements in or rela
ITITLE OF INVENTION: agents
FILE REFERENCE: REF/Klaveness/206
CURRENT APPLICATION NUMBER: US/08/959,206A
CURRENT FILLING DATE: 1997-10-24
NUMBER OF SEQ ID NOS: 27
SEQ ID NO 7
LENGTH: 10
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                                                                                                                                      APPLICANT: Nycomed
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Best Local
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LENGTH: 10
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Patent No. 626491
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 71.9%;
Best Local Similarity 80.0%;
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILE REFERENCE: REF/KLaveness/993
CURRENT APPLICATION NUMBER: US/08/958,993A
CURRENT FILING DATE: 1997-10-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Nycomed Imaging AS
TITLE OF INVENTION: Improvements in
TITLE OF INVENTION: agents
FILE REFERENCE: REF/Klaveness/903
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                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: AMIDATION
                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION:
NAME/KEY: MOD RES
LOCATION: (10)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: MOD_RES
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Biotinylated-Gly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Description of Artificial Sequence:Biotinylated fibrin-anti-polymerant peptide
                                                                                                       Improvements in or relating to diagnostic/therapeutic
                                                                                                                                                                                                                                                             10
                                                                                                                                                                                                                                                                                                                                  71.9%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
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Pred. No. 0
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Query Match
Best Local Similarity
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                                                                             ; MOLECULE TYPE: protein US-08-749-816-2
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US-08-749-816-2
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                                                                                                                                                                                                                                                                         COMPUTER: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/749,816
FILING DATE: 15-NOV-1996
CLASSIFICATION: 536
ATTORNEY/AGENT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
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                                                                                                                                   TELEFAX: 215-875-8394
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 336 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: MOD_RES
LOCATION: (1)
OTHER INFORMATION: Biotin-Gly
NAME/KEY: MOD RES
LOCATION: (10)
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                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: Welser, Gerard J.
REGISTRATION NUMBER: 19,763
REFERENCE/DOCKET NUMBER: 98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Romey, Georges
APPLICANT: Barhanin, Jacques
TITLE OF INVENTION: FAMILY OF MAMMALIAN POTASSIUM CHANNELS,
TITLE OF INVENTION: THEIR CLONING AND THEIR USE ESPECIALLY FOR THE SCREENING
TITLE OF INVENTION: OF DRUGS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
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                                                                                                          TOPOLOGY:
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Guillemare, Eric
Fink, Michel
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                Conservative
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                                                                                                                                                                                                      215-875-8383
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                             64.9%;
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80.0%;
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                             Score 37; DE
Pred. No. 13;
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Pred. No. 0.054;
                Mismatches
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                                            Length 336;
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                Indels
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US-09-144-914-2
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                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: SEIDEL, CHRISTOPH
APPLICANT: SEIDEL, PETER
APPLICANT: VON DER ELTZ, HERBERT
TITLE OF INVENTION: SYNTHETIC STANDAF
NUMBER OF SEQUENCES: 11
CORRESSONDENCE ADDRESS:
ADDRESSEE: NIKAIDO, MARMELSTEIN, MADDRESSEE: METROPOLITAN SQUARE, 655 15
STREET: 330-G ST.LOBBY
CITY: WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-08-762-695-11
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SEQ ID NO 2
LENGTH: 336
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: FAMILY OF MAMMALIAN POTASSIUM CHANNELS, THEIR CLONING TITLE OF INVENTION: AND THEIR USE, ESPECIALLY FOR THE SCREENING OF DRUGS FILE REFERENCE: 989.6705CIP CURRENT APPLICATION NUMBER: US/09/144,914

CURRENT FILING DATE: 1998-09-01

EARLIER APPLICATION NUMBER: 08/749,816

EARLIER FILING DATE: 1996-01-15

EARLIER FILING DATE: 1996-01-5,234

EARLIER APPLICATION NUMBER: 60/095,234

EARLIER FILING DATE: 1996-02-08

EARLIER FILING DATE: 1996-02-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Fink, Michel APPLICANT: Lazdunski, Michel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Duprat, APPLICANT: Lesage,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: TWIK-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Homo sapiens FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT
                                                                                                                                        ZIP: 20005-5701
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                             SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/762,695
                   APPLICATION NUMBER: US/0:
FILING DATE: 12-DEC-1996
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                             STATE: D.C
COUNTRY: U
  APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12 RLVERHRSA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3 RVVERHOSA 11
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77.8%;
08/325,589
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Pred. No. 13;
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FILING DATE: 20-OCT-PRIOR APPLICATION DATA:

APPLICATION NUMBER: DE P FILING DATE: 20-OCT-1993

19-OCT-1994

43 35

798.9

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Query Match
Best Local Similarity 1
Totales 7; Conservati
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US-08-592-500-40
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US-08-762-695-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 40, App. Sequence 40, App. 6005089
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: Lanza, Francois
APPLICANT: Phillips, David R.
APPLICANT: Cazenave, Jean-Pierre
TITLE OF INVENTION: Platelet Glycoprotein V Gene and Uses
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 11 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (202)638-4810
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:

APPLICATION NUMBER: DE P 44 2

FILING DATE: 15-JUN-1994

ATTORNEY/AGENT INFORNATION:

NAME: NOLAN, SHARON L.

REGISTRATION NUMBER: 36,335

REFERENCE/DOCKET NUMBER: P161

TELECOMMUNICATION INFORMATION:

TELEPHONE: (202)638-5000
                                                                                                          NAME: Dow, Karen B.
REGISTRATION NUMBER: 29,684
REFERENCE/DOCKET NUMBER: 12418-28
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 326-2400
TELEPHONE: (415) 326-2400
                                                                                                                                                                                                                                                                                                                       FILING DATE:

CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 08/089,455
FILING DATE: 09-JUL-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FORM:
COMPUTER: IBM FORM:
COMPUTER: Floppy disk
COMPUTER:
COMPU
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Palo Alto
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/592,500
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TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE:
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FILING DATE: 20-MAY-1994
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    Townsend and Townsend Khourie and Crew
379 Lytton Avenue

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                                                       5) 320
/ 326-2422
- NO: 40:
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 7;
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                                                                                                                   US-08-592-500-40
                                                     Query Match 63.2
Best Local Similarity 100.
Matches 7; Conservative
                                                                                                                                                                                                                                                                                       HYPOTHETICAL: NO FEATURE
                                                                                                                                                                                                                                                                                                            TYPE: amino acid
TOPOLOGY: unknown
MOLECULE TYPE: pept
                                                                                                                           LOCATION: 5..7
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                       LOCATION: 1..11
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                             NAME/KEY:
                                                                                                                                                                                                        OTHER INFORMATION:
                                                                                                                                                                                                                                                      NAME/KEY: Peptide LOCATION: 1..11
                          1 GPRVVER 7
                                                                                                                                                            Region 5..7
                                                                63.2%; Score 36; DB 3; 100.0%; Pred. No. 0.55;
                                                                                                                              /note= "Amino acid residues
identical to GPV."
                                                                                                                                                                                                        cleavage site."
                                                                                                                                                                                                                /note= "Amino acid sequence of the human fibrinogen (Fg) A-alpha 2 chain thrombin
                                                       0,
                                                       Mismatches
                                                    0; Indels
                                                                               Length 11;
                                                    0,
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0

Search completed: June 27, 2003, 17:19:03 Job time : 13.1 secs

0,

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OM protein -
protein search, using sw model
                                                          Copyright
                                                        GenCore version 5.1.6 (c) 1993 - 2003 Compugen Ltd.
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June 27, 2003, 17:15:16 ; Search time 11.7 Seconds (without alignments) 73.950 Million cell updates/sec

Title: Perfect score: Sequence: US-10-019-439-2. 51 1 GHRPLDKKR 9

Run on:

Scoring table: BLOSUM62 Gapop 10.0 , Gapext 0.5

Searched: 283224 segs, 96134422 residues

Total number of hits satisfying chosen parameters: 283224

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 su

summaries

PIR_73:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*

Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14	13	12	11	10	9	00	7	o	ຫ	4	w	N	_	No.	Result
33	33	33	33	33	33	u	ω ω	34	34	34	34	34	34	34	35	ω	36	36	36	37	38	39	39	39	39	43	51		Score	
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485	464	436	424	403	250	162	155	862	795	736	480	419	402	399	571	225	1490	201	195	275	463	673	231	231	231	468	491	31	Length	
N	2	N	N	N	N	N	ν	N	N	N	N	N	N	N	N	N	2	N	2	2	N	N	N	N	μ	-	ا	N	BB	
F70464	H90340	AB3419	D57742	T20727	T46417	S66764	856258	B36786	A84608	T05137	G75072	T03675	E97398	AE2616	AG3174	A75550	T47840	C95399	C95286	T04480	A38463	T40817	D85814	D90966	F64960	FGBOB	FGHUB	A05297	·ID	,
hypothetical prote	etical p	protei	ا چ	_	Ľ	ר	_		hypothetical prote	protein kinase hom	proly1-tRNA synthe	cyclin 2 - rice	ın.	ical	acetolactate synth	MutT/nudix family	ев		Φ	acyl-CoA oxidase h		77		ical prot	membrane protein y		fibrinogen beta ch	en b	Description	

RESULT 2
FORHUB
fibrinogen beta chain precursor [validated] - human
fibrinogen beta chain precursor [validated] - human
N;Alternate names: coagulation factor I
N;Contains: fibrinopeptide B
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Jate: 24-Apr-1984 #sequence revision 31-Mar-1993 #text change 08-Dec-2000
C;Jate: 24-Apr-1986; A90469; B90469; I37389; A94433; A90437; A94309; G54223; A03121; B3
R;Chung, D.W; Harris, J.E.; Davie, E.W.
Adv. Exp. Med. Biol. 281, 39-48, 1990

A; Title: Nucleotide sequences of the three genes coding for human fibrinogen.

45	44	43	42	41	40	39	38	37	36	35	34	ω ω	32	31	30
32	32	32	32	32	32	32	32	32	32	32	33	ω W	3 3	ω G	33
62.7	62.7	62.7	62.7	62.7	62.7	62.7	62.7	62.7	62.7	62.7	64.7	64.7	64.7	64.7	64.7
271	262	222	215	199	183	168	152	147	142	127	1058	888	682	570	501
2	ผ	N	N	μ.	ν	N	N	N	N	2	N	N	N	N	N
D70591	B72333	H83387	S23432	D33830	AH0804	S78110	н90358	H69180	S39708	G95291	D82654	D84650	T18847	G95326	H84727
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hypothetical prote	bacteriocin - Ther	probable hydrolase	pyroglutamyl-pepti	cation efflux syst	conserved hypothet	thi protein - Rhiz	hypothetical prote	conserved hypothet	ywdC protein - Bac	hypothetical prote	ankyrin-like prote	probable potassium	hypothetical prote	probable acetolact	probable inositol

ALIGNMENTS

Qy 1 GHRPLDKKR 9 Db 20 GHRPLDKKR 28
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A; Molecule type: protein
A; Residues: 164-174 < KUN>
A; Residues: 164-174 < KUN>
A; Residues: 164-174 < KUN-
A; Rote: identification of tryptic peptides from high-densi R; Henschen, A; Lottspeich, F.; Kehl, M.; Southan, C.
Ann. N. Y. Acad. Sci. 408, 28-43, 1983
A; Fitle: Covalent structure of fibrinogen.
A; Reference number: A90037; MUID:83254370; PMID:6575689
A; Contents: annotation; review, disulfide bonds
R; Garddlund, B, Hessel, B.; Marguerie, G.; Murano, G.; Blc
Bur. J. Biochem. 77, 595-610, 1977
A; Fitle: Primary structure of human fibrinogen. Character;
A; Reference number: A91249; MUID:77245999; PMID:891553
                                                                                                                                                                                                            A;Title: The structures of fibrinogen and
                                                                                                                                                                                                                                                                                                           A; Contents: annotation; disulfide bonds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AGCESSION..., AG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Thromb. Res. 8, 639-658, 1976
A;Title: Disulfide bridges in NH-2-terminal part of human fibrinogen
A;Reference number: A94309; MUID:76225080; PMID:936108
A;Contents: disulfide bonds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Molecule type: protein
A; Residues: 31-144, 'QF',147-231,'D',233-330,'E',332-491
R; Blomback, B.; Hessel, B.; Hogg, D.
                                                                                                                                                                                                                                                  ;Doolittle, R.F.; Takagi, T.; Watt, K.; Boon Regulatory Proteolytic Enzymes and Their
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Title: Amino acid sequence of the beta chain of human
A;Reference number: A90437; MUID:79124640; PMID:420779
A;Accession: A90437
                                                                                                                                                   Contents: annotation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Blochemistry 18,
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R;Huber, P.; Dalmon, J.; Courtois, G.; Laurent, M.; Assouline, Z.; Marguerie, G.
Nucleic Acids Res. 15, 1615-1625, 1997
A;Title: Characterization of the 5'-flanking region for the human fibrinogen beta gene.
A;Reference number: 137389; MUID:87146483; PMID:3029722
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cross-references: EMBL:X05018; NID:g31400; PIDN:CAA28674.1; PID:gHenschen, A.; Lottspeich, F.; Southan, C.; Topfer-Petersen, E. n Protides of the Biological Fluids, Proc. 28th Colloq., Peeters, Title: Human fibringen: sequence, sulfur bridges, glycosylation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Reference number: A43568; MUID:91344740; PMID:2102623
A;Accession: B43568
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Molecule type: protein
Residues: 31-137,'QS',140-144,'QF',147-491
Watt, K.W.K.; Takagi, T.; Doolittle, R.F.
ochemistry 18, 68-76, 1979
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contents: carbohydrate binding
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Molecule type: DNA
Residues: 1-38 <HUB>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ACCESSION: A94433
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Residues: 1-38 <CH1>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Molecule type: DNA Residues: 9-191,'P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Que,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ue, B.G.; Rixon, M.W.; Mace Jr., 3244-3250, 1983
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             B.G
                                                                                                                                                   disulfide
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9; MUID:83283433; PMID:6688356
                                                                                                                                                                                                                                                                             K.; Bouma III, H.;
                                                                                                                                                                                                                                                                                                                                                                                                                                            G.; Murano, G.; Blomback,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          from high-density
Southan, C.
                                                                                                                                                                                                                                                                                                                                                         Characterization of disulfide-containing
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Magnusson
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structural
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A;Residues: 1-4 <BLO>
R;Sjoquist, J.; Blomback,
Ark. Kemi 16, 425-436, 196
                                                                                                      A; Molecule type:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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N/Contains: fibrinopeptide B
C/Species: Bos primigenius taurus (cattle)
C/Date: 29-Jul-1981 #sequence revision 29-Jul-1981 #text_change 13-Sep-1996
C/Accession: A03122; B03117; B37507; A37513; S02443
R/Blomback, B.; Doolittle, R.F.
Acta Chem. Scand. 17, 1816-1819, 1963
A/Title: The sequence of amino acids at the N-terminal end of bovine fibring A;Reference number: A03122
A/Boscession: A03122
                                                                                                                                                                                                                                                                                                                                  fibrinogen beta chain - bovine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Pathway: blood coagulation
C;Superfamily: fibrinogen beta chain; fibrinogen beta/gamma homology; fibrinogen disulf
C;Keywords: blood coagulation; coiled coil; glycoprotein; liver; plasma; pyroglutamic a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   R;Kirschbaum, N.E.; Budzynski, A.Z.
J. Biol. Chem. 265, 13669-13676, 1990
A;Title: A unique proteolytic fragment of human fibrinogen containing the A;Reference number: A37117; MVID:90337977; PMID:2143188
A;Contents: annotation; hementin cleavage site
C;Comment: The conversion of fibrinogen to fibrin is triggered by thrombi: tration sites responsible for the formation of the soft clot.
C;Comment: The conversion of fibrinogen to fibrin is triggered by thrombi: C;Comment: The soft clot is converted into the hard clot by factor XIIIa ger) and between alpha chains (weaker) of different monomers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Description: fibrinogen cleaved
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Cross-references: GDB:119130; OMIM:134830
A;Map position: 4q28-4q28
A;Introns: 38/3; 102/3; 164/1; 240/1; 278/1; 320/1; 415/2
C;Complex: The fibrinogen molecule is a hexamer containing tins are contained in the core. Two three-chain coiled coils
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ;106/Disulfide bonds: interchain (to alpha-68) #status experimental;110/Disulfide bonds: interchain (to gamma-45) #status experimental;223/Disulfide bonds: interchain (to alpha-184) #status experimental;227/Disulfide bonds: interchain (to gamma-161) #status experimental;231-316;241-270,424-437/Disulfide bonds: #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ann. N. Y. Acad. Sci. 408, 449-456, 1983
A;Title: Cloning of fibrinogen genes and their cDNA.
A;Reference number: A90038; MUID:83254384; PMID:6575700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ;238-487/Domain: fibrinogen beta/gamma homology <FBG>
;31/Modified site: pyrrolidone carboxylic acid (GIn) (in mature form)
;44-45/Cleavage site: Arg-Gly (thrombin) #status experimental
;95/Disulfide bonds: interchain (to alpha-55) #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ;45-47/Region: polymerization site
;99-228/Domain: fibrinogen disulfide ring homology <FDR>
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;1-30/Domain: (or 4-30 or 15-30) signal sequence #status predicted
;31-491/Product: fibrinogen beta chain #status experimental <MAT>
;31-44/Product: fibrinopeptide B #status experimental <APT>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    from the distal domain nodes.
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Y. Acad. Sci. 408, 449-456, 1983
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
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                                                            protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   carbohydrate (Asn) (covalent)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   53
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Wallen,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 51;
Pred. No.
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                                                                                                                  end of bovine fibrinopeptide
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s emerge from this
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thrombin, which
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R;Medved, L.V.; Platonova, T.N.; Litvinovich, S.V.; Lukinova, N.I.
FEBS Lett. 232, 56-60, 1988
A;Title: The cleavage of beta-chain in bovine fibrinogen D(H) fragment (95
A;Reference number: S02443; MUID:88211875; PMID:2966748
A;Accession: S02443
A;Accession: S02443
A;Residues: 373-374 (MED)
C;Comment: Thrombin cleaves the bond between Arg-21 and Gly-22 to release
C;Comment: Thrombin cleaves the bond between Arg-21 and Gly-22 to release
                                                                 C;Keywords: transmembrane protein F;48-64/Domain: transmembrane #status F;135-151/Domain: transmembrane #statt
                                                                                                                                                                                                                                                                                                                                                                                                                                                    C;Accession: F64960
R;Blattner, F.R.; Plunkett III,
.A.; Rose, D.J.; Mau, B.; Shao,
                                                                                                                                                                                                       A;Cross-references: GB:AE000288; GB:U00096; NID:g2367124; PIDN:AAC75028.1; PID:g1788273
A;Experimental source: strain K-12, substrain MG1655
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Species: Escherichia coli
C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 01-Mar-2002
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A;Title: Amino acid sequences of portions of the alpha A;Reference number: A37507; MUID:79164394; PMID:434821
A;Accession: B37507
                                                                                                                                                                                                                                                                            A; Molecule type: DNA
A; Residues: 1-231 < BLAT>
                                                                                                                                                                                                                                                                                                                                                     Science 277, 1453-1462, 1997
A;Title: The complete genome sequence of Escherichia coli K-12.
A;Reference number: A64720; MUID:97426617; PMID:9278503
A;Accession: F64960
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              7:215-464/Domain: fibrinogen beta/gamma homology 7FBGs
7:1/Modified site: pyrrolidone carboxylic acid (GIn) #status experimental
7:6/Binding site: sulfate (Tyr) (covalent) #status experimental
7:21-22/Cleavage site: Arg-Gly (thrombin) #status experimental
7:371/Binding site: carbohydrate (Asn) (covalent) #status predicted
7:372-373/Cleavage site: Arg-Thr (plasmin) #status experimental
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Comment: Fibrinogen is a hexamer containing two sets of three nonidentical charce superfamily: fibrinogen beta chain; fibrinogen beta/gamma homology; fibrinogen ketwords: blood coagulation; glycoprotein; plasma; pyroglutamic acid; sulfoprot; 76-205/Domain: fibrinogen disulfide ring homology FDR>
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Query Match
Best Local Similarity
                                                                                                                                                Superfamily: conserved hypothetical protein AF0994
                                                                                                                                                                             Gene:
                                                                                                                                                                                                  Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Species: Escherichia coli
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Best Local
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                                                                            transmembrane #status
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76.5%;
87.5%;
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  Score
Pred.
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  39;
                                                                         predicted <TM2>
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                       BB
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                                                                                                      <TM1 >
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                       Length 231;
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                                                                                                                                                                                                                                                                                                                              shown
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 chains
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iller, L.; Grotbeck, E.J.; Davis, N.W.; Lim, A.; Dimalanta, Nature 409, 529-533, 2001
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                                                                                                                  C; Superfamily:
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C;Superfamily:
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A;Residues: 1-231 <HAY>
A;Cross-references: GB:BA000007; PIDN:BAB36123.1; PID:g13362168; GSPDB:GN00154
A;Experimental source: strain O157:H7, substrain RIMD 0509952
A;Experimental source: strain O157:H7, substrain RIMD 0509952
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Reference number: A99629; A; Accession: D90966
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          R;Hayashi, T.; Makino,
gasawara, N.; Yasunaga,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         R;Hayashi,
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                                                                                                                                                                                             A;Molecule type: DNA
A;Residues: 1-231 <STO>
                                                                                                                                                                                                                         A;Status: preliminary
                                                                                                                                                                                                                                                     A;Title: Genome sequence of enterohemorrhagic Escherichia A;Reference number: A85480; MUID:21074935; PMID:11206551
                                                                                                                                                                                                                                                                                                                                                                            hypothetical protein yedJ [imported] - Escherichia coli (strain O157:H7, substrain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Title: Complete genome sequence of enterohemorrhagic Escherichia coli O157:H7
A;Reference number: A99629; MUID:21156231; PMID:11258796
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA Res. 8,
                                                                                                                                                                                                                                         A; Accession:
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                                                         Best Loc
Matches
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                                                                                                                                                 Genetics:
                                                                                                                                                                 ;Experimental source:
                                                                                                                                                                                                                                                                                                                                             Date: 16-Feb-2001 #sequence_revision 16-Feb-2001 #text_change 14-Sep-2001
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Escherichia coli
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                                                           Similarity 7; Conser
                                                                                                                                                                                                                                         D85814
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HRPLDDKR
                            HRPLDKKR 9
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Yasunaga, T.; Kuhara,
                                                                                                                     conserved
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                                                           Conservative
                                                                                                                                                               GB:AE005174; NID:g12516071; PIDN:AAG56976.1; ce: strain O157:H7, substrain EDL933
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                                                                       76.5%;
                                                                                                                  hypothetical protein AF0994
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Pred. No.
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Shiba, T.;
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                                                                                      DB 2; Length 231;
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lanta, E.;
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M.; Shinagawa,
                                                                                                                                                                                                                                                                      0157:H7.
                                                                                                                                                                                                                                                                                                    J.D.; Rose, Potamousis,
                                                                                                                                                                               GSPDB:GN00145; UWGP:Z3
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zinc finger protein - fission y c;Species: Schizosaccharomyces C;Date: 03-Dec-1999 #sequence_r C;Accession: T40817

protein - fission yeast (Schizosaccharomyces pombe)

#sequence_revision 03-Dec-1999

#text_change

02-Sep-2000

RESULT

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C;Species: 'Gaille 1991 #sequence_...
C;Date: 31-Jul-1991 #sequence_...
C;Date: 31-Jul-1991 #sequence_...
C;Accession: A38463
R;Weissbach, L.; Oddoux, C.; Procyk, R.; Grieninger, G.
Biochemistry 30, 3290-3294, 1991
A;Title: The beta chain of chicken fibrinogen contains an information of the contains of the contains of the contains and contai
                                                                                                                                                                                                                                                  RiGrossi, M.; Gulli, M.; Stanca, A.M.; Cattivelli, A; Title: Characterization of two barley genes that A; Reference number: Z15371
A; Accession: T04480
                                                                                                                   A;Cross-references: EMBL:X84055; NID:g642243; PIDN:CAA58874.1; A;Experimental source: cv. Arda, leaf
                                                                                                                                                                                          A; Molecule type: mRNA
A; Residues: 1-275 < GRO>
                                                                                                                                                                                                                                       A; Status: preliminary; translated
                                                                                                                                                                                                                                                                                                                                                                                                        acyl-CoA oxidase homolog - barley (fragment)
C;Species: Hordeum vulgare (barley)
C;Date: 23-Apr-1999 #sequence_revision 23-Apr-1999 #text_change 08-Oct-1999
C;Accession: T04480
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A,Residus: 1-463 <WEIN
A,Cross-references: GB:M58514; NID:g211779; PIDN:AAA48770.1; PID:g211780
C;Superfamily: fibrinogen beta chain; fibrinogen beta/gamma homology; filt
F;73-202/Domain: fibrinogen disulfide ring homology FDR>
F.712-460/Domain: fibrinogen beta/gamma homology <FBG>
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Best Local Sin
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A;Residues: 1-673 <BEC>
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A;Experimental source: strain 972h-; clone pl p8B7
C;Genetics:
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submitted to the EMBL Data
A;Reference number: Z21949
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ibrinogen beta chain - chicken (fragment)
;Species: Gallus gallus (chicken)
;Date: 31-711-1991 #sequence_revision 31-Jul-1991 #text_change
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        Similarity
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75.0%;
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66.7%;
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Library, October 1998
                                                                                                                                                                                                                                                                                       barley genes that respond rapidly to dehydration stress
  Score
Pred.
                                                                                                                                                                                                                                       from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 38;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pred. No. 13;
2; Mismatches
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Pred. No. 12;
1; Mismatches
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  37;
No.
DB
12;
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                Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 673
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                                                                                                                                    PID:g642244
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A; Residues: 1-201 < KUR>
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A; Cross-references: GB.AE006469; PIDN:AAK64853.1; PID:914523267; GSPDB:GN00165 A; Experimental source: strain 1021, megaplasmid pSymA R; Galibert, F.; Finan, T.M.; Long, S.R.; Puhler, A.; Abola, P.; Ampe, F.; Barlt, D.; Cchain, P.; Cowie, A.; Davis, R.W.; Dreano, S.; Federspiel, N.A.; Fish Science 293, 668-672, 2001
A; Authors: Kahn, D.; Kahn, M.L.; Kalman, S.; Keating, D.H.; Kiss, E.; Komp, C. A; Title: The composite genome of the legume symbiont Sinorhizobium meliloti. A; Reference number: A96039; MUID:21368234; pMID:11474104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R;Barnett, M.J.; Fisher, R.F.; Jones, T.; Komp, C.; Abola, A.P.; Barloy-Hubler, F.; Bow.; Kalman, S.; Keating, D.H.; Palm, C.; Peck, M.C.; Surzycki, R.; Wells, D.H.; Yeh, K.C.; Proc. Natl. Acad. Sci. U.S.A. 98, 9883-9888, 2001
A;Title: Nucleotide sequence and predicted functions of the entire Sinorhizobium melilo A;Reference number: A95262; MUID:21396509; PMID:11481432
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Molecule type: DNA
A; Residues: 1-195 < KUR>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     probable transposase [imported] - Sinorhizobium meliloti (strain 1021) magaplasmid
C;Species: Sinorhizobium meliloti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Status: preliminary
                                                                                                                                                                                                       Gene: SMa0360
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Parte: 24-Aug-2001 #sequence_revision 24-Aug-2001 #text_change
                                                                                                                                                                                  plasmid
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                                                                                         Similarity 7; Conserv
                                      GHRPLDKKR 9
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54
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77.8%;
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Pred. No. 13;
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                                                                             Mismatches
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                                                                                                                      2;
                                                                                                                 Length 195;
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S.; Federspiel, N.A.; Fisher, R.F
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probable transcription regulator [imported] - Sinorhizobium meliloti (strain C;Species: Sinorhizobium meliloti C;Date: 24-Aug-2001 #sequence_revision 24-Aug-2001 #text_change 30-Sep-2001 C;Accession: C95399 2001 #sequence_revision 24-Aug-2001 #text_change 30-Sep-2001

1021)

maga

R;Barnett, M.J.; Fisher, R.F.; Jones, T.; Komp, C.; Abola, A.P.; Barloy-Hubler, F.; Bow.; Kalman, S.; Keating, D.H.; Palm, C.; Peck, M.C.; Surzycki, R.; Wells, D.H.; Yeh, K.C. Proc. Natl. Acad. Sci. U.S.A. 98, 9883-9888, 2001
A;Title: Nucleotide sequence and predicted functions of the entire Sinorhizobium melilo A;Reference number: A95262; MUID:21396509; pMID:11481432

Barloy-Hubler, Fisher, R.F.

A;Cross-references: GB.AE006469; PIDN:AAK65757.1; PID:g14524255; GSPDB:GN00165 A;Experimental source: strain 1021, megaplasmid pSymA R;Galibert, F; Finan, T.M.; Long, S.R.; Puhler, A.; Abola, P.; Ampe, F.; Barle Pela, D.; Chain, P.; Cowie, A.; Davis, R.W.; Dreano, S.; Federspiel, N.A.; Fish Science 293, 668-672, 2001
A;Authors: Kahn, D.; Kahn, M.L.; Kalman, S.; Keating, D.H.; Kiss, B.; Komp, C. A;Title: The composite genome of the legume symbiont Sinorhizobium meliloti. A;Reference number: A96039; MUID:21368234; PMID:11474104

C.; Lelaure

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RESULT 13
A75550
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                                                                                                                                                                                                                                                                       R; White, O.; Eisen, J.A.; Heidelberg, J.F.; Hickey, I.M.; Shen, M.; Vamathevan, J.J.; Lam, P.; McDonald, S.; Smith, H.O.; Venter, J.C.; Fraser, C.M.
Science 286, 1571-1577, 1999
A; Title: Genome sequence of the radioresistant bacte.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Map position: 3
A;Introns: 9/1; 795/3; 824/3; 1036/3; 1075/3; 1174/1; 1245/3; 1347/3; 1369/1; 1449/1
A;Note: T209.140
C;Superfamily: human multidrug resistance protein cMOAT2; ATP-binding cassette homolo
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A;Cross-references: EMBL:AL138658
A;Experimental source: cultivar Columbia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Gene: SMa2008
A;Genome: plasmid
                                                                                                                                                                                                                                               A;Accession: A75550
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A;Accession: T47840
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submitted to the Protein Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        multi resistance protein homolog - Arabidopsis thaliana
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                                                                                                                                                                                                                                                           ;Title: Genome sequence of the radioresistant bacterium ;Reference number: A75250; MUID:20036896; PMID:10567266
                                                                                                                                                                   ;Cross-references: GB:AE001881;
;Experimental source: strain R1
                                                                                                                                                                                                                  ;Molecule type: DNA
                                                                           Query Match
Best Local
                                                                                                                                                                                                Residues: 1-225 <WHI>
                                                                                                                                                                                                                                Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                Species: Deinococcus radiodurans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Alternate names: protein T209.140
Species: Arabidopsis thaliana (mouse-ear cress)
                                                              Matches
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GHRELDKR 185
                            GHRPLDKK 8
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                                                           Conservative
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100.0%;
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75.0%;
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lequence Database,
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Pred. No.
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Pred. No.
                                                                        Score 35; DB
Pred. No. 24;
                                                            Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20-Apr-2000 #text_change 21-Jul-2000
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l, L.; Utterback,
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T.; Zalewski,
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47 HRPVDQER

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Search completed: June 27, Job time : 12.7 secs

2003, 17:18:36

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C; Superfamily: o
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A; Title: The Genome of t)
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R;Wood, D.W.; Setubal, J.C.; Kaul, R.; Monks, erage, G.; Gillet, W.; Grant, C.; Guenthner, Karp, P.; Romero, P.; Zhang, S. Science 294, 2317-2323, 2001
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R;Title: The Genome of the Natural Genetic Engineer Agrobacterium tumefaciens C58.
A;Reference number: AB2577; PMID:11743193
                                                                                                                                                                                                                                                                                                                                            A;Title: The Genome of the Natural Genetic Engineer Agrobacterium tumefaciens A;Reference number: AB2577; PMID:11743193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       hypothetical protein hemN [imported] - Agrobacterium tumefaciens (strain C58, C;Species: Agrobacterium tumefaciens C;Date: 11-Jan-2002 #sequence_revision 11-Jan-2002 #text_change 01-Feb-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Cross-references: GB:AE008687; PIDN:AAL45813.1; PID:g17743551; GSPDB:GN00188
A;Experimental source: strain C58 (Dupont)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R;Wood, D.W.; Setubal, J.C.; Kaul, R.; Monks, D.; Chen, L.; Wood, G.E.; Chen, Y.; Woo, erage, G.; Gillet, W.; Grant, C.; Guenthner, D.; Kutyavin, T.; Levy, R.; Li, M.; McClel; Karp, P.; Romero, P.; Zhang, S.
Science 294, 2317-2323, 2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C;Species: Agrobacterium tumefaciens
C;Date: 11-Jan-2002 #sequence_revision 11-Jan-2002 #text_change 01-Feb-2002
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                                                                                                                                                                                                                                                                           , Molecule type: DNA
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A;Accession: AG3174
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                                                                                                                                                                                                               ;Cross-references: GB:AE008688; PIDN:AAL41347.1; ;Experimental source: strain C58 (Dupont)
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Protides of the biological fluids,
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"Human fibrinogen: sequence,
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Nickerson D.A.;
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cinogen, thrombosis,
num Press, New York
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hemistry 18:68-76(1979).
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sequences of
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(Rel. 26, Last sequence update)
(Rel. 41, Last annotation update)
beta chain precursor (Contains: Fibrinopeptide
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Hessel B., Hogg D.;
idges in NH2-terminal
                                                                    ; PubMed=3029722;
J., Courtois G.;
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15:1615-1625(1987)
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Yoshida N., Wada H
Asakura S., Shirak
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Asakura S., Shirakawa S.,
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Spraggon G., Everse
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MEDLINE=98292395; PubMed=9628725;
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MEDLINE=77245999; Pul
Gaardlund B., Hessel
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SEQUENCE OF 31-44.

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PubMed=1634610;
kate F., Lord S.T.,
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Liu C.Y., Koehn J.A., Morgan F.J.;
"Characterization of fibrinogen New York 1.
fibrinogen with a deletion of B beta(9-72)
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DISEASE: DEFECTS IN FGB ARE A CAUSE OF THROMBOPHILIA.

DISEASE: DEFECTS IN FGB ARE A CAUSE OF THROMBOPHILIA.

MISCELLANEOUS: CONVERSION OF FIBRINOGEN TO FIBRIN IS TRIGGERED BY
THROMBIN, WHICH CLEAVES FIBRINOPEPTIDES A AND E FROM ALPHA & BETA
CHAINS, AND THUS EXPOSES THE N-TERMINAL POLYMERIZATION SITES
RESPONSIBLE FOR THE FORWATION OF THE SOFT CLOT. THE SOFT CLOT IS
CONVERTED INTO THE HARD CLOT BY FACTOR XIIIA WHICH CATALYZES THE
EPSILON-(GAMMA-CLUTAMYL)LYSINE CROSS-LINKING BETWEEN GAMMA CHAINS
(STRONGER) AND BETWEEN ALPHA CHAINS (WEAKER) OF DIFFERENT
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L; J00130; AAA98115.1; J0INED.
L; J00132; AAA98116.1; -...
L; J00133; AAA98216.1; -...
L; AF388026; AAK62470.1; -...
L; AK638036; AAA5245.1; -...
L; M66876; AAA52445.1; J0INED.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SUBUNIT: HEXAMER CONTAINING 2 SETS OF 3 NONIDENTICAL CHAINS (ALPHA, BETA AND GAMMA), LINKED TO EACH OTHER BY DISULFIDE BONDS THE AMINO ENDS OF ALL CHAINS ARE CONTAINED IN THE CENTRAL WODULE ARE 2 THREE-CHAIN COLLED COLLS, WHICH CONNECT THE CENTRAL NODULE ARE 2 THREE-CHAIN COLLED COLLS, WHICH CONNECT THE CENTRAL NODULE TO THE DISTAL NODULES CONTAINING THE DISTAL DOMAINS. EXTENDING FAR PERIPHERALLY ARE THE LONG CARBOXYL
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31ol. Chem. 260:4390-4396(1985).
FUNCTION: FIBRINGGEN HAS A DOUBLE
POLYMERIZE INTO FIBRIN AND ACTING
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.A., Morgan F.J.;
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Blomback B., Dooli
"The sequence of a
fibrinopeptide B."
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Proc. Natl. Acad. Sci. U.S.A. 7
Proc. Natl. Acad. Sci. U.S.A. 7
-i- FUNCTION: FIBRINOGEN HAS A
POLYMERIZE INTO FIBRIN AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Martinelli R.A., Inglis A.S., Rubira Hurrell J.G.R., Leach S.J., Scheraga "Amino acid sequences of portions of bovine fibrinogen.";
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Mammalia; Eutheria;
Bovidae; Bovinae; B
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MEDLINE=81199473; PubMed=6262803;
Chung D.W., Rixon M.W., McGillivray
"Characterization of a cDNA clone cc
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Sjoquist J., Blo
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                                                                                                                                                                                                                                                                                                                                                                                                                               SUBUNIT: HEXAMER CONTAINING 2 SETS OF 3 NONIDENTICAL CHAINS (ALPHA, BETA AND GAMMA), LINKED TO EACH OTHER BY DISULETIDE BONDS. MISCELLANGOUS: CONVERSION OF FIBRINGEN TO FIBRIN IS TRIGGERED BY THROMBIN, WHICH CLEAVES FIBRINOPEPTIDES A AND B FROM ALPHA & BETA CHAINS, AND THUS EXPOSES THE N-TERMINAL POLYMERIZATION SITES RESPONSIBLE FOR THE FORMATION OF THE SOFT CLOT: THE SOFT CLOT IS CONVERTED INTO THE HARD CLOT BY FACTOR XIIIA WHICH CATALYZES THE EBSILON-(GAMMA-GLUTAMYL)LYSINE CROSS-LINKING BETWEEN GAMMA CHAINS (STRONGER) AND BETWEEN ALPHA CHAINS (WEAKER) OF DIFFERENT
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email to license@isb-sib.ch).
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e of amino
                                                                                                                                        fibrinogen_C; 1.
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Last annotation update)
precursor (Contains: Fibrinopeptide
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ovine fibrinopeptides
                                                  C DOMAIN; 1.

Platelet; Glycoprotein;

FIBRINOPEPTIDE B.

FIBRINOGEN BETA CHAIN
PYRROLIDONE (SULFATION. CLEAVAGE (BY
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Scheraga H.A.;
tions of the al
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RESULT
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STRAIN-K12 / MG1655,

( MEDLINE-97426617; PubMed-9278503;
A Blattner F.R., Plunkett G. III, Bloch C.A.,
A Riley M., Collado-Vides J., Glasner J.D., R.
Davis N.W., Kirkpatrick H.A., Go
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Itoh T., Aiba H., Baba T., Fujita K., Hayashi K., Inada T.,
Isono K., Kasai H., Kimura S., Kitakawa M., Kitagawa M.,
Makino K., Miki T., Mizobuchi K., Mori H., Mori T., Motomur
Saito N., Sampei G., Nashimoto H., Nishio Y., Oshima T.,
Takeda J., Takemoto K., Wada C., Yamamoto Y., Horiuchi T.;
"A 460-kb DNA sequence of the Escherichia coli K-12 genome
DNA Res. 3:379-392(1996).
                      "Detection of new genes in a bacterial three gene classes.";
Nucleic Acids Res. 23:3554-3562(1995).
                                                                                    Danchin
                                                                                             MEDLINE=96032851;
Borodovsky M., McI
                                                                                                                                                                                        STRAIN=K12;
MEDLINE=89345179;
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P46144; P76333;
01-NOV-1995 (Rel. 32, Created)
01-NOV-1997 (Rel. 35, Last sequence up
15-JUN-2002 (Rel. 41, Last annotation
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                                                                                                                                              ., Gerwin N., Fritz H.-J.;
tide sequence of the dcm locus of
Acids Res. 17:5844-5844(1989).
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structure.";
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Mammalia; Eutheria; Rodentia;
MCBI_TaxID=10116;
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Rattus norvegicus (Rat).
Univervota: Metazoa; Chordata;
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EMBL; X13330; -; Yead.

EMBL; X13330; -; Yead.

EcoGene; EG12710; Yead.

InterPro; IPR002819; HD.

InterPro; IPR003607; ME_Pplase_HDc.

Pfam; PF01966; HD; 1.

SMART; SM00471; HDc; 1.

Tengthetical protein; Complete proteome.

Tength
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HRPLDDKR 168
                                                                                                    Conservative
                                                                                                                          76.5%;
87.5%;
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                                                                                                                     Score 39;
Pred. No.
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P14480;
Ol-JAN-1990 (Rel. 13, Created)
Ol-OCT-1996 (Rel. 34, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation updat
Fibrinogen beta chain precursor [Contains: STANDARD; :ation update)
[Contains: Fibrinopeptide

Courtney M.A., Bunce L.A., "Cloning of the complete co STRAIN=Sprague-Dawley; TISSUE=Liver; MEDLINE=95143386; PubMed=7841303; ., Neroni L.A., Simpson-Haidaris p. coding sequence of rat fibrinogen conservation of fibrin beta 15-42 Craniata; Vert Sciurognathi; Vertebrata; Euteleostomi; Muridae; Murinae; Rattus B beta primary

MEDINE-84194000, PubMed=6232608;
MEDINE-84194000, PubMed=6232608;
Fowlkes D.M., Mullis N.T., Comeau C.M., Crabtree G.R.;
Potential basis for regulation of the coordinately ex
fibrinogen genes: homology in the 5' flanking regions.
Proc. Natl. Acad. Sci. U.S.A. 81:2313-2316(1984) Blood Coagul. Fibrinolysis 5:487-496(1994). SEQUENCE OF 1-26 FROM N.A expressed

fibrinogen: evolutionary conservation 3'-untranslated sequences."; Gene 79:151-158(1989) SEQUENCE OF 183-479 FROM N.A. MEDLINE=89378771; PubMed=2673932; Eastman E.M., Gilula N.B.; "Cloning and characterization of a "Cloning and Characterization" Blomback B., Blomback M., Grondahl N., "Studies on fibrinopeptides from mamm. Acta Chem. Scand. 19:1789-1791(1965). Grondahl N.J.;
es from mammals."; CDNA of translated for the B beta chain

N.A.

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RESULT 6
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EMBL; M27220; AAA41160.1; -.
EMBL; K01336; AAA98625.1; -.
EMBL; M35602; AAA41159.1; -.
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                                                                                                                                                                                                                                                                                                                                                                                                                             InterPro; IPR002181; Fibrinogen_C.
Pfam; PF00147; fibrinogen_C; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SUBUNIT: HEART AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                . Celī Res. 169:47-56(1987).
FUNCTION: FIBRINOGEN HAS A
POLYMERIZE INTO FIBRIN AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SUBUNIT: HEXAMER CONTAINING 2 SETS OF 3 NONIDENTICAL CHAINS (ALPHA, BETA AND GAMMA), LINKED TO EACH OTHER BY DISULFIDE BOND MISCELLANEOUS: CONVERSION OF FIBRINGEN TO FIBRIN IS TRIGGERED THROMBIN, WHICH CLEAVES FIBRINOPEPTIDES A AND B FROM ALPHA & BE CHAINS, AND THUS EXPOSES THE N-TERMINAL POLYMERIZATION SITES RESPONSIBLE FOR THE FORMATION OF THE SOFT CLOT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              A05299;
PE0010;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SIMILARITY: CONTAINS 1 FIBRINOGEN C-TERMINAL DOMAIN
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                                                      CHICK
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S
                                                                                                                                                                                                                                                                                                                                                                                               coagulation; Plasma;
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E; PS00514; FIBRIN_AG
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                                                                                                                                                                Similarity
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beta chain
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33; PubMed=3817019;
tti A.-M., Taroux P.,
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                                                      STANDARD;
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258
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Last sequence update)
Last annotation update)
precursor [Contains: Fibrinopeptide
                                 Created)
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LSI -> ILS (IN REF. 3)
L -> Q (IN REF. 5).
S -> T (IN REF. 5).
S -> K (IN REF. 5).
R -> K (IN REF. 5).
                                                                                                                                                     4;
                                                                                                                                                                                                                                                                                                                                                                                                        C DOMAIN; 1.
                                                                                                                                                  Score 39; DB
Pred. No. 2.8;
4; Mismatches
                                                                                                                                                                                                                                                                                                                                                               FIBRINOPEPTIDE
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                                                      463
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  (See http://www.isb-sib.ch/announce/
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2.8;
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Gallus gallu
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SITE
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SEQUENCE
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NON TER 1
PEPTIDE <1
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MEDLINE=91182745; PubMed=2009266;
Weissbach L., Oddoux C., Procyk R
                                                                                                                                                                                                                                                                                                                         PIR; A38463; A38463.
                                                                                                                                                                                                                                                                                                                                                                              modified
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-!- FUNCTION: FIBRINOGEN HAS A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cleavage site.";
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                                                                                                                                                                                                                                                                       PROSITE;
                                                                                                                                                                                                                                                                                         Pfam; PF00147; fibrinogen_C;
                                                                                                                                                                                                                                                                                                                 HSSP;
                                                                                                                                                                                                                                                                                                                                    EMBL; M58514; AAA48770.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "The beta chain of chicken
                                                                                                                                                                                                                                                                                                     InterPro; IPR002181;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                SUBUNIT: HEXAMER CONTAINING 2 SETS OF 3 NONIDENTICAL CHAINS (ALPHA, BETA AND GAMMA), LINKED TO EACH OTHER BY DISULFIDE BONDS. MISCELLANEOUS: CONVERSION OF FIBRINGEN TO FIBRIN IS TRIGGERED BY THROMBIN, WHICH CLEAVES FIBRINGPEPTIDES A AND B FROM ALPHA & BETA CHAINS, AND THUS EXPOSES THE N-TERMINAL POLYMERIZATION SITES RESPONSIBLE FOR THE FORMATION OF THE SOFT CLOT IS CONVERTED INTO THE HARD CLOT BY FACTOR XILLA WHICH CATALYZES THE EPSILON-(GAMMA-GLUTAMYL)LYSINE CROSS-LINKING BETWEEN GAMMA CHAINS (STRONGER) AND BETWEEN ALPHA CHAINS (WEAKER) OF DIFFERENT
                                                                                                                                                                                                                                                                                                                                                                                      s SWISS-PROT entry is copyright. It is produ
ween the Swiss Institute of Bioinformatics
European Bioinformatics Institute. There a
by non-profit institutions as long as
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SUBUNIT: HEXAMER CONTAINING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FUNCTION: FIBRINOGEN HAS A POLYMERIZE INTO FIBRIN AND
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19
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                                        Similarity 6; Conserv
                                                                                                                                                                                                                                                                     PS00514;
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HRPLDKRQ 26
               HRPLDKKR
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80
84
197
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                                        Conservative
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463
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nogen_C; 1.
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BY SIMILARITY.
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N-2044CD49BA79EC7B (
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CLEAVAGE (BY THROWN
FIBRINOPEPTIDE B).
INTERCHAIN (WITH A)
INTERCHAIN (WITH G)
INTERCHAIN (WITH G)
INTERCHAIN (WITH G)
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Pred. No.
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                                        ore 38; DB
med. No. 4.3;
Mismatches
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                                                                                                                                                                                                                                                           Glycoprotein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FUNCTION: YIELDING MONOMERS AS A COFACTOR IN PLATELET
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                                                             Length 463;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     an atypical thrombin
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                                                                                   CRC64;
                                         Indels
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RESULT 7 MRAW_ZYMMO

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MEDLINE=98442695; PubMed=9771752;
Buxbaum J.D., Choi E.K., Luo Y., Lilliehook C., Crowley A.C.
Merriam D.E., Wasco W.;
"Calsenilin: a calcium-binding protein that interacts with
presenilins and regulates the levels of a presenilin fragme
Nat. Med. 4:1177-1181(1998).
                                                                                                                                                                                                                                                                                                                                                                                                                             Q9YZW7; Q9UJ84; Q9UJ85;
16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Calsenilin (DRB-antagonist modulator) (DREAM) (Ky
Protein 3) (A-type potassium channel modulatory pr
CSEN OR DREAM)
CSEN OR DREAM (Human)
                                                          MEDLINE=99176420; PubMed=10078534; Carrion A.M., Link W.A., Ledo F.,
                                                                                                                                      SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on it modified and this statement is not removed. Usage by and for commercia entities requires a license agreement (See http://www.isb-sib.ch/announce or send an email to license@isb-sib.ch).
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STRAIN=ATCC 31821 / ZM4 / CP4;
STRAIN=ATCC 31821 / ZM4 / CP4;
SUM H.W., Kang H.S.;
Submitted (AUG-1999) to the EMBL/GenBank/DDBJ databases.
Submitted (AUG-1999) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OPERQ; OPRIMO; Created)
15-JUN-2002 (Rel. 41, Created)
15-JUN-2002 (Rel. 41, Last sequence update)
15-JUN-2002 (Rel. 41, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
5-adenosyl-methyltransferase mraw (EC 2.1.1.-).
MRAW OR ZM110RF5 OR ZM16ORF1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Zymomonas mobilis.
Bacteria; Proteobacteria; alpha subdivision; Sphingomonadaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=542;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FUNCTION: Exhibits a S-adenosyl-dependent activity (By similarity).
SIMILARITY: BELONGS TO THE MRAW FAMILY.
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75.0%;
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                                                       Mellstrom B.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (See http://www.isb-sib.ch/announce/
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                                                     Naranjo J.R.;
                                                                                                                                                                                                                                                                Crowley A.C.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Kv channel-interacting
protein 3) (KChIP3).
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Best Local
Oryza sativa (Rice)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
snermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
                                                                                   Q40671;
15-DEC-1998 (Rel. 37, Created)
15-DEC-1998 (Rel. 37, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
G2/mitotic-specific cyclin 2 (B-like cyclin)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nuclear protein.
CA BIND 175
CA-BIND 223
CONFLICT 182
CONFLICT 207
SEQUENCE 256 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SMART; SM00054; EFh; 3.
PROSITE; PS00018; EF_HAND;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      InterPro; iPR002048; EF-hand.
InterPro; IPR001125; Recoverin.
Pfam; PF0036; efhand; 3.
PRINTS; PR00450; RECOVERIN.
ProDom; PD00012; EF-hand; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This SWISS-PROT entry is copyright. It is produced through a c between the Swiss Institute of Bioinformatics and the EMBL the European Bioinformatics Institute. There are no restrict use. by non-profit institutions as long as its content is modified and this statement is not removed. Usage by and fo entitles requires a license agreement (See http://www.isb-sib.or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Genew; HGNC:15523; CSEN MIM; 604662; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Transcription regulation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.

MEDLINE=20140134; PubMed=10676964;

An W.F., Bowlby M.R., Betty M., Cao J.; Ling H.-P., Mendoza G.,

Hinson J.W., Mattson K.I., Strassle B.W., Trimmer J.S., Rhodes K.J.

"Modulation of A-type potassium channels by a family of calcium
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SUBGUNIT: BINDS TO DNA AS A HOMOMULTIMER. ASSOCIATES WITH C-TERMINUS OF PSEN1 AND PSEN2. ASSOCIATES WITH KCN1.
SUBCELLULAR LOCATION: CYTOPLASMIC. ALSO MEMBRANE-BOUND. NUCLEAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INDUCTION: AFFINITY FOR DNA IS SIMILARITY: CONTAINS 2 EF-HAND
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398:80-84(1999).
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EF-HAND 2 (BY SIMILARITY)
I -> V (IN REF 2)
R -> Q (IN REF 2)
4W; 635C3EDF8B91E1C5 CRC64;
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MEDLINE=95261415; PubMed=7742859;
Sauter M., Mekhedov S.L., Kende H.;
"Gibberellin promotes histone H1 kinase activity and the expression of cdc2 and cyclin genes during the induction of rapid growth in deepwater rice internodes.";
                                                                                                                                                                                                                                                                                                                                                                                                                      Pfam; PF00134; cyclin; 1.
Pfam; PF02984; cyclin C; 1.
SMART; SM00385; CYCLIN; 2.
PROSITE; PS00292; CYCLINS; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
               SEQUENCE FROM
STRAIN=Auburn
                                                    Ictalurid herpesvirus 1 (Channel catfish virus) (CCV).
Viruses; dsDNA viruses, no RNA stage; Herpesviridae;
ICtalurid Herpes-like viruses.
NCBI_TaxID=10401;
                                                                                                                                                                                                                     HSVI1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "Isolation and characterization of a cDNA encoding a mitotic cyclin the CycB2 type from rice.";
(In) Plant Gene Register PGR97-001.
-!- FUNCTION: ESSENTIAL FOR THE CONTROL OF THE CELL CYCLE AT THE G2/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EMBL; X82036; CAA57556.1; -. HSSP; P30274; IVIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sauter M.;
MEDLINE=92087490; PubMed=1727613;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            InterPro; IPR004366; Cyclin.
InterPro; IPR004367; Cyclin_Cterm
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DEVELOPMENTAL STAGE: ACCUMULATES STEADILY DURING ABRUPTLY DESTROYED AT MITOSIS.
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SUBUNIT: INTERACTS WITH THE CDC2 PROTEIN KINASE TO FORM A
SERINE/THREONINE KINASE HOLOENZYME COMPLEX ALSO KNOWN AS
MATURATION PROMOTING FACTOR (MPF). THE CYCLIN SUBUNIT IMPARTS
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5; Conserv
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419 AA; 47572 MW; A8774B56BD839A5B CRC64;
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(Rel.
(Rel.
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                                                                                                                                                                        24, Created)
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62.5%;
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Last annotation updat
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                                                                                                                                                                                                                                                                          use by modified
PROSITE; PS00232; CADHERIN_1; 17.
PROSITE; PS50268; CADHERIN_2; 27.
Cell adhesion; Glycoprotein; Tran
Signal; Multigene family.
POTE
SIGNAL
1 23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15-JUN-2002
15-JUN-2002
15-JUN-2002
                                                                                                             EMBL, AB053447; BAB61904.1; -.
InterPro; IPR002126; Cadherin.
Pfam; PF00028; cadherin; 27.
SMART; SM00112; CA; 26.
                                                                                                                                                                                                                                                                                                                                                                    This
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=21481446; PubMed=11597768;
Nakajima D., Nakayama M., Kikuno R., Hirosawa
"Identification of three novel non-classical c
comprehensive analysis of large cDNAs.";
Brain Res. Mol. Brain Res. 94:85-95(2001).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL; M75136; AAA88182.1; -. EMBL; M75136; AAA88104.1; -. PIR; B36786; B36786.
                                                                                                                                                                                                                             entities requires a license agreement (S or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                            between
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Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cadherin 23 precursor (Otocadherin). CDH23 OR KIAA1774.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Rattus norvegicus (Rat).
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by non-profit institutions as long as
lified and this statement is not removed. I
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FUNCTION: Cadherins are calcium dependent cell adhesion proteins.

They preferentially interact with themselves in a homophilic

They preferentially interact with the homophilic
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CADHERIN 23.
EXTRACELLULAR (POTENTIAL).
POTENTIAL.
CYTOPLASMIC (POTENTIAL).

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ID CADN H
AC 09H251
DT 15-JUN
DT 15-JUN
DT 15-JUN
DT 15-JUN
DT 15-JUN
DT CADNER
OS Homo s
OC Bukary
OC Mammal
RP SEQUEN
RP H-1496
RP N-1350
RP H-1496
RP N-1361
RP L-2380
RP MEDLIN
RA BOLZ H
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RA Nothwa
RA BIT Causes
RN [2]
RY Wayne
RA Griffi
RT Usher
RT Usher
RT Griffi
RT Gena Griffi
RT Gena
RA Nakajina D., Nakayama M., Kikuno R., Hirosawa M., Nagase T., Ohara O., RT "Identification of three novel non-classical cadherin genes through RT comprehensive analysis of large cDNAs."

RI Brain Res. Mol. Brain Res. 94:85-95 (2001).

CC -!- FUNCTION: Cadherins are calcium dependent cell adhesion proteins. CC They preferentially interact with themselves in a homophilic bundle organizer perhaps by cross-linking the stereocilia (By companie in connecting cells. Cadherin 23 may function as hair country).

CC similarity).

CC -!- SUBCELIULAR LOCATION: Type I membrane protein (By similarity).

CC and 4; are produced by alternative splicing.

CC and 4; are produced by alternative splicing.

CC TISSUE SPECIFICITY: Particularly strong expression in the retina.

CC OLISEASE: Defects in CDH23 are the cause of Usher syndrome type 1D characterized by congenital profound sensorineural hearing loss; constant vestibular dysfunction and prepubertal onset of retinitis retinitis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              밁
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Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bork J.M., Peters L.M., Riazuddin S., Bernstein S.L., Ahmed Z.M., Ness S.L., Polomeno R., Ramesh A., Schloss M., Srisailpathy C.R.S. Wayne S., Bellman S., Desmukh D., Ahmed Z., Khan S.N., Kaloustian V.M.D., Li X.C., Lalwani A., Riazuddin S., Bitner-Glindzicz M., Nance W.E., Liu X.-Z., Wistow G., Smith R.J.: "Usher syndrome ID and nonsyndromic autosomal recessive deafness DFNB12 are caused by allelic mutations of the novel cadherin-like
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A., ALTERNATIVE SPLICING, VARIANTS USHID M-1281 DEL;
H-1496 AND Q-1746, AND VARIANTS C-3; A-490; N-496; T-1222; C-1349;
N-1351; A-1575; S-1671; I-1675; Q-1804; S-1999; K-2044; Q-2358;
L-2380; Q-2588 AND L-3125;
MEDLINE=20578759; PubMed=11138009;
BOIZ H., Von Brederlow B., Ramirez A., Bryda E.C., Kutsche K.,
Nothwang H.G., Seeliger M., Del C.-Salcedo Cabrera M.,
Vila Caballero M., Pelaez Molina O., Gal A., Kubisch C.;
"Mutation of CDH23, encoding a new member of the cadherin gene famil."
"Mutation of CDH23, encoding a new member of the cadherin gene famil."
Nat. Genet. 27:108-112(2001)."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE=21481446; PubMed=11597768;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE OF 2173-3354 FROM N.A. (ISOFORM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE OF 803-3354 FROM N.A., ALTERNATIVE SPLICING, AND VARIANTS DFNB12 N-990; N-2045; N-2202; N-2950; C-2956 AND T-3059.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
NCBI TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ISSUE=Brain;
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Catarrhini; Hominidae
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Pred. No.
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SMART; SM00112; CA; Zb.

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PROSITE; PS00232; CADHERIN 1; 17.

PROSITE; PS50268; CADHERIN 2; 27.

Cell adhesion; Glycoprotein; Transmembrane; Calcium-binding; Repeat; Cell adhesion; Glycoprotein; Transmembrane; Calcium-binding; Repeat; Signal; Multigene family; Alternative splicing; Vision; Polymorphism; Signal; Multigene family; Alternative splicing; Vision; Polymorphism; Disease mutation; Usher syndrome; Deafiness; Retinitis pigmentosa.

Signal 1 23 POTENTIAL.

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EMBL; AY010111; AAG27034.2;
EMBL; AB053445; BAB61902.1;
HSSP; P15116; 1NCJ.
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MIM; 601386;
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MIM; 605516; -.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cause of combined dea DISEASE: Defects in C autosomal recessive d SIMILARITY: Contains
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    s requires a license agreement (S an email to license@isb-sib.ch).
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    27 cadherin domains.
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/FTId=VAR_012171.
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RP SEQUENCE FROM N.A.

RC STRAIN=C59BL/6J; TISSUB=Testis;
RX MEDLIN=21085660; PubMed=1121781;
RA ARANACTORDAMA A., Shibata K., Voshino M., Itoh M., Ishii Y.,
RA Arakawal J., Shinagawa A., Shibata K., Konno H., Adachi J., Pukuda S.,
RA Arakawa T., Hara A., Fukunishi Y., Konno H., Kondo S., Yamanaka I.,
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RA Aizawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I.,
RA Aizawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I.,
RA Saito T., Okazaki Y., Gojobori T., Bono H., Kasukawa T., Saito R.,
RA Kidota K., Matsuda H.A., Ashburner M., Batalov S., Casavant T.,
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RA Kuehl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Schriml L.M., Staubli F., Suzuki R., Tomita M., Wagner L., Washio T.,
RA Saki K., Okido T., Furuno M., Aono H., Baldarelli R., Batshi G.,
RA Saki K., Okido T., Furuno M., Aono H., Baldarelli R., Washio T.,
RA Blake J., Boffelli D., Bojunga N., Carminci P., de Bonaldo M.F.,
RA Blake J., Bottoni L., Mashima J., Mazarelli H., Gariboldi M.,
RA Brownstein M.J., Bult C., Fletcher C., Fujita M., Gee N.H.,
RA Brownstein M.J., Bult C., Fletcher C., Fujita M., Gee N.H.,
RA Brownstein M.J., Batoka J., Mazarelli J., Mombaerts P.,
RA Gustincich S., Hill D., Hofmann M., Hume D.A., Kamiya M., Lee N.H.,
RA Brownstein M., Sato K., Schoenbach C., Seya T., Shibata Y., Storch K.-F.,
RA Wynshaw-Boris A., Yoshida K., Hasegawa Y., Kawaji H., Kohtsuki S.,
RT "Bincritonal Brownstein M., Schoenbach C., Whittaker C., Wilming L.,
RT "Bincritonal Brownstein M., Schoenbach C., Seya T., Shibata Y., Storch K.-F.,
RT "Bincritonal Brownstein M., Schoenbach C., Whittaker C., Wilming L.,
RT "Bincritonal Brownstein M., Schoenbach C., Seya T., Shibata Y., Storch K.-F.,
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Q99PF4; Q99NH1; Q99NN9;

15-UUN-2002 (Rel. 41, Created)

15-UUN-2002 (Rel. 41, Last sequence updated)

15-UUN-2002 (Rel. 41, Last annotation updated)

Cadherin 23 precursor (Otocadherin)
           alleles of cadherin
Gene 281:31-41(2001)
                      pi Palma F., Pellegrino R., Noben-Trauth "Genomic structure, alternative splice for alleles of cacherin 23 (Cdh23).";
                                                                      MEDLINE=21623040; PubMed=11750125;
                                                                                                                                                               SEQUENCE FROM N.A. Fu Y., Wang Q., Roe Submitted (SEP-2000)
                                                                                                                    GENOMIC ORGANIZATION,
2718-ASN-GLU-PRO-2720
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE=20578758; PubMed=11138008;
Di Palma F., Holme R.H., Bryda E.C., Belyantseva T.A., Pellegrino Kachar B., Steel K.P., Noben-Trauth K.;
"Mutations in Cdh23, encoding a new type of cadherin, cause stereocilia disorganization in waltzer, the mouse model for Usher syndrome type 1D.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Wilson S.M., Householder D.B., Coppola V., Tessarollo L., Fritzsch B., Lee E.-C., Goss D., Carlson G.A., Copeland N.G., Jenkins N.A., "Mutations in Cdh23 cause nonsyndromic hearing loss in waltzer mice.", Genomics 74:228-233(2001).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=21280917; PubMed=11386759;
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Mammalia; Eutheria;
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GLU-PRO-2720 DEL, AND VARIANTS
ARG-1236; VAL-2025; VAL-2026;
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Sciurognathi; Muridae; Murinae; Mus
                                                                                            SPLICING, VARIANT WALZER RIANTS PRO-5; VAL-229; LYS-, 2026; THR-2217; HIS-2222;
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PROSITE; PS50268; CADHERIN 2; 27.
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SUBCELLULAR LOCATION: Type I membrane protein (By similarity).

ALTERNATIVE PRODUCTS: 2 isoforms; 1 (shown here) and 2; are

produced by alternative splicing.

TISSUE SPECIFICITY: In adult animals relatively high levels of expression are found in testis, skeletal muscle, hear, eye and the sensory half cells of the inner ear.

DISBASE: Defects in CDH23 are the cause of waltzer (v) phenotype. Waltzer mice are characterized by deafness and vestibular dysfunction due to degeneration of the neuroepithelium within the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SIMILARITY: Contains 27 cadherin domains.
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AC079818; -; NOT_ANNOTATED_CDS.
AC079819; -; NOT_ANNOTATED_CDS.
AC079082; -; NOT_ANNOTATED_CDS.
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                        EMBL; D50617; BAA09242.1;
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Saccharomycetales;
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Brown A.E., Jackson P
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Okinaka R.T., Cloud
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SIMILARITY: TO S.POMBE S
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SEQUENCE FROM N.A.

STRAIN=5288c / AB972;

MEDLINE=95400292; PubMed=7670463;

Murakami Y., Naitou M., Hagiwara H., Shi
Sasanuma S.-I., Sasanuma M., Tsuchiya Y.
Yamazaki M., Tashiro H., Eki T.;
"Analysis of the nucleotide sequence of
Saccharomyces cerevisiae.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Saccharomyces cerevisiae (Baker's yeast).
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is SWISS-PROT entry is copyright. It is produced through a collaboration tween the Swiss Institute of Bioinformatics and the EMBL outstation - between the Swiss Institute. There are no restrictions on its e Buropean Bioinformatics Institute. There are no restrictions on its by non-profit institutions as long as its content is in no way diffied and this statement is not removed. Usage by and for commercial titles requires a license agreement (See http://www.isb-sib.ch/announce/send an email to license@isb-sib.ch).
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pXO2-72.
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protein in NIC96-MPR1
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Pred. No. 6.6;
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a Y., Soeda
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KW Hypothetical protein.
FT DOMAIN 93 103 POLY-SER.
SQ SEQUENCE 155 AA; 18187 MW; E191184DCD5EF8A1 CRC64;

Query Match
Best Local Similarity 75.0%; Pred. No. 13;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 2 HRPLDKKR 9
Db 123 HRKLEKKR 130

Search completed: June 27, 2003, 17:17:09

Search completed: June 27, 2003, 17:17:09

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Result
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sp_rodent:*
sp_virus:*
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Q9VWQ3
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Q92XY9
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OBww77 homo sapien
Q8xb89 escherichia
Q94271 schizosacch
Q46583 desulfovibr
Q93ju5 pseudomonas
Q43476 hordeum vul
Q95rf0 drosophila
Q9vwq5 rhizobium m
Q92xy9 rhizobium m
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Q98x90 homo sapien
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RESULT 2
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QBWW77;
Q1-MAR-2002 (TrEMBLrel. 20, Created)
Q1-MAR-2002 (TrEMBLrel. 20, Last sequence update)
Q1-JUN-2002 (TrEMBLrel. 21, Last annotation update)
Hypothetical 46.9 kDa protein.
Hypothetical 46.9 kDa protein.
Homo sapiens (Human).
                                                                                                                                                                       Strausberg R.;
Submitted (JAN-2002) to the EMBL/GenBank/DDBJ databases.
EMBL; BC02762; AAH20762.1; -.
InterPro; IPR002181; Fibringen C.
InterPro; IPR001019; Gprotein_alpha.
Pfam; PF00147; fibringen_C; 1.
Pfam; PF00503; G-alpha; 1.
SMART; SM00186; FBG; 1.
SMART; SM00186; FBG; 1.
                                                                                                                                                      Hypothetical SEQUENCE 41
                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                  Q8WW77
                                                                                                                                                                                                                                                                                       TISSUE=LIVER;
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411 AA; 40
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094271;
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01-JUN-2002
Beck A., Reinhardt R, Lyne M., Submitted (CCT-1998) to the EMBMBI; AL032684; CAA21808.1; -. InterPro; IPR001841; Znf_ring.
SMART; SM00184; RING; 1.
                                                                                                                                                                                                   Schizosaccharomyces pombe (Fission yeast).
Eukaryota; Fungi; Ascomycota; Schizosaccharomycetes;
Schizosaccharomycetales; Schizosaccharomycetaceae;
                                                                                    STRAIN=972H-;
Beck A., Rein
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01-MAR-2002
01-MAR-2002
01-JUN-2002
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Hayashi T., Makino K., Ohnishi M., Kurokawa K., Ishii Hayashi T., Ohtsubo E., Nakayama K., Murata T., Tanaki Iida T., Takami H., Honda T., Sasakawa C., Ogasawarak Kuhara S., Shiba T., Hattori M., Shinagawa H.; "Complete genome sequence of enterchemorrhagic Escher O157:H7 and genomic comparison with a laboratory stranscript of the comparison with a laboratory with a laboratory 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Perna N.T., Plunkett G. III, Burland V., Mau B., Glasner J.D., Rose D.J., Mayhew G.F., Evans P.S., Gregor J., Kirkpatrick H.A., Posfai G., Hackett J., Klink S., Boutin A., Shao Y., Miller L., Grotbeck E.J., Davis N.W., Lim A., Dimalanta E.T., Potamousis K., Apodaca J., Anantharaman T.S., Lin J., Yen G., Schwartz D.C., Welch R.A., Blattner F.R.,
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EMBL; AP002559; BAB36123.1; -.
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Escherichia coli O157:H7.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=21074935; PubMed=11206551;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       interPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=83334;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    161
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PF01966; HD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity 7; Conser
                                                                                                                         FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HRPLDKKR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    proteome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HRPLDDKR 168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       IPR002819; HD.
IPR003607; ME_Pplase_HDc
1966; HD; 1.
                                                                                                                                                                                                                                                                                                              (TrEMBLrel.) (TrEMBLrel.) (TrEMBLrel.)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25835 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  76.5%;
87.5%;
                                                                                                                                                                                                                                                                                                           10,
10,
21,
                                       M., Rajandream M.A
e EMBL/GenBank/DDBJ
; -.
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                                                                                                                                                                                                                                                                                                       Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                                                                                   Created)
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Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 39;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Created)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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                                                                              М.А.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 update)
                                                          databases
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                                                                            Barrell B.G.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ishii
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Tobe T.,
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Best Local S
Matches 6
                                                                              ProDom; PD006099; Hemery
SMART; SM00304; HAMP; 1.
SMART; SM00283; MA; 1.
                                                                                                                           Pfam;
                                                                                                                                                                                                                                                                                                                                                                                                                                          01-NOV-1996
01-NOV-1996
01-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PROSITE; PSO
Zinc-finger.
SEQUENCE 6
                                                                                                           Pfam; PF01814; Hemerythrin; Pfam; PF00015; MCPsignal; 1
                                                                                                                                                                                                                            Deckers H.M., Voordouw G.;
"The dcr gene family of Desulfovibrio: im
of dcrH and phylogenetic comparison with
Antonie Van Leeuwenhoek 70:21-29(1996).
                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
STRAIN=HILDENBOROUGH;
                                                                                                                                                                       EMBL; U30319; AAB50497.1;
                                                                                                                                                                                  Submitted
                                                                                                                                                                                                   STRAIN=HILDENBOROUGH;
                                                                                                                                                                                                              SEQUENCE FROM N.A
                                                                                                                                                                                                                                                                                                                                                                                              Desulfovibrio vulgaris (st
Bacteria; Proteobacteria;
Desulfovibrio.
                                                                                                                                                                                                                                                                                                                                                                                                                          DCRH.
                                                                                                                                      InterPro;
                                                                                                                                                      InterPro;
                                                                                                                                                                 HSSP; P02942; 1QU7
                                                                                                                                                                                           Voordouw G.
                                                                                                                                                                                                                                                                MEDLINE=96433451; PubMed=8836438;
                                                                                                                                                                                                                                                                                                     Hildenborough genome.";
J. Bacteriol. 176:351,358(1994)
                                                                                                                                                                                                                                                                                                                     "Identification of a large family proteins in an ordered library of
                                                                                                                                                                                                                                                                                                                                        MEDLINE=94117369; PubMed=8288529; Deckers H.M., Voordouw G.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Q46583
                                                                                                                                                                                                                                                                                                                                                            STRAIN=HILDENBOROUGH;
                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE OF 605-727 FROM
                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=882;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Q46583;
                                                                                                                                               nterPro;
513
                                                                                                                          PF00672; HAMP;
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                                  Similarity 7; Conserv
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HRPLDKAR
                  HRPLDKKR
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                                                                                                                                  IPR003660; HAMP.
IPR002063; Hemerythrin.
                                                                                                                                                   IPR004089; Chmtaxis_transd
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                                                                                                                                                                                  (JUN-1995) to
                                                                        959
                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                        (TrEMBLrel. (TrEMBLrel. (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  673 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                      AA;
                                                                                                          MCPsignal;
520
                  9
                                                                                                Hemerythrin;
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                                                                      104729 MW;
                                            76.5%;
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66.7%;
                                                                                                                                                                                                                                                                                                                                                                                                     (strain Hildenborough)
ia; delta subdivision;
                                                                                                                                                                                the EMBL/GenBank/DDBJ databases
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                                  Score 39; DB
Pred. No. 32;
0; Mismatches
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Last sequence update)
Last annotation updat
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Pred.
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                                                                    450550E840954D42 CRC64;
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e Desulfovibrio vulo
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32;
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23;
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                                  Indels
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RESULT 5 Q93JUS ID Q93JUS

PRELIMINARY;

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RESULT
Q95RF0
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Q43476
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Matches 6
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Best Local
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Submitted (JUN-2001) to the EMBL/Ger
EMBL; AJ313205; CAC42170.1; -.
InterPro; IPR004830; LRV:
InterPro; IPR004831; Unk_DeinSyn.
Pfam; PF01816; LRV; 5.
ProDom; PD038723; Unk_DeinSyn; 1.
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Q43476;
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SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-NOV-1996 (TrEMBLrel. 01, 01-NOV-1996 (TrEMBLrel. 01, 01-UN-2002 (TrEMBLrel. 21,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-NOV-1996
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01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation updat
Hypochetical 27.9 kDa protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRAIN=CV. ARDA; TISSUE=C
Grossi M., Gulli M., Star
"Characterization of two
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Viridiplantae; Stro
Spermatophyta; Magnoliophyta;
Triticeae; Hordeum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mammalian acyl CoA oxidase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        dehydration stress.";
Plant Sci. 105:71-80(1995).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Desnoues N., Lin M., Elmerich "Organisation of nif genes in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRAIN=CHINA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            InterPro; IPR001552; Acyl-CoA_dh.
InterPro; IPR005202; GRAS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=4513;
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                                                                                                                                 225
                                                                                                                                                                                                                                                                                                                                                                                                                                                           PF00441; Acyl-CoA_dh; 1.
PF03514; GRAS; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         X84055; CAA58874.1;
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                                                                                                                                 GHRPLRRRR 233
                                                                                                                                                                                              GHRPLDKKKR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GHRPLNRK 187
                                                                                                                                                                                                                                                                                                                                                                                             275
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                                                                                                                                                                                                                                                                                                                                                                                             AA;
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M., Stanca A.M., C
n of two barley gen
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                                                                                                                                                                                                                                                                                                                                                                                             30689 MW;
                                                                                                                                                                                                                                                                                              72.5%;
66.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  74.5%;
75.0%;
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yta; Liliopsida; Poales; Poaceae; Pooideae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    , Last sequence update)
, Last annotation update)
homologous (Fragment).
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Pred.
                                                                                                                                                                                                                                                                                              Score 37;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C.;
Pseudomonas stutzeri A15,
                                                                                                                                                                                                                                                                                                                                                                                             332811FD05827472 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                               Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         38; DB
No. 13;
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                                                                                                                                                                                                                                                                                                                             В
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                                                                                                                                                                                                                                                                                                                         Length 275;
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A Adams M.D., Celniker S.E., Li P.W., Hoskins R.A., Galle R.F.,
A Amanatides P.G., Scherer S.E., Li P.W., Hoskins R.A., Galle R.F.,
A George R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N.,
RA George R.A., Lewis S.E., Yandell M.D., Zhang Q., Chen L.X.,
RA Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X.,
RA Brandon R.C., Rogers Y.-H.C., Blazej R.G., Champe M., Pfeiffer B.D.,
RA Brandon R.C., Baxter E.G., Helt G., Nelson C.R., Miklos G.L.G.,
RA Abril J.F., Agbayani A., An H.-J., Andrews Pfannkoch C., Baldwin D.,
RA Ballew R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley E.M.,
RA Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S.,
RA Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P.,
RA Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA de Pablos B., Delcher A., Deng Z., Mays A.D., Dew I., Dietz S.M.,
RA Dodson K., Doup L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P.,
RA Fosler C., Gabrielian A.E., Garg N.S., Gebbart W.M., Glasser K.,
RA Glodek A., Gong F., Gorrell J.H., Gu Z., Guan P., Harris M.,
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Best Local S
Matches 6
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Q9VWQ6;
01-MAY-2000
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Q95RF0;
01-DEC-2001
01-DEC-2001
01-JUN-2002
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CG6961.
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CG18259 protein.
CG18259.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRAIN=BERKELEY;
MEDLINE=20196006; PubMed=10731132;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Drosophila melanogaster (Fruit fly).
Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;
Pterygota; Neoptera; Endopterygota; Diptera; Brachycera; Musc
Ephydroidea; Drosophilidae; Drosophila.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FlyBase; FBgn0030959; CG6961.
InterPro; IPR000504; RNA_rec_mot.
Pfam; PF00076; rrm; 1.
PRINTS; PR01656; VACCYTOTOXIN.
PROSTITE; PS50102; RRM; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Stapleton M., Brokstein P., Hong L., Agbayani A., Carlson J., Champe M., Chavez C., Dorsett V., Farfan D., Frise E., George Gonzalez M., Guarin H., Li P., Liao G., Miranda A., Mungall C. Nunco J., Pacleb J., Paragas V., Park S., Phouanenavong S., Wa V. C., Levis S.E., Rubin G.M., Celniker S.; Submitted (OCT-2001) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-MAY-2000
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) (TrEMBLrel. 13,
2 (TrEMBLrel. 21,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRELIMINARY;
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75.0%;
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Last annotation updat
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Last sequence update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 37; DB
Pred. No. 40;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    44C8C8DA263899B4 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Brachycera; Muscomorpha;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Brachycera; Muscomorpha;
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K.A.,

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AX MEDLINE=20196006; PubMed=10731132;

A Adams M.D. Celniker S.E., Holt R.A., Evans C.A., Gocayne J.D.,

A Adams M.D., Celniker S.E., Holt R.A., Hoskins R.A., Galle R.F.,

A Amanatides P.G., Scherer S.E., Li P.W., Hoskins R.A., Galle R.F.,

A George R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N.,

A Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X.,

RA Brandon R.C., Rogers Y.-H.C., Blazej R.G., Champe M., Pfeiffer B.D.,

RA Man K.H., Doyle C., Baxter E.G., Helt G., Nelson C.R., Miklos G.L.G.,

RA Abril J.F., Agbayani A., An H.-J., Andrews Pfannkoch C., Baldwin D.,

RA Ballew R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley E.M.,

RA Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S.,

RA Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P.,

RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,

RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,

RA Cherny J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,

RA Dodson K., Doup L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P.,

RA Durbin K.J., Evangelista C.C., Ferriara C., Ferriera S., Fleischmann W.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RA Harris N.L., Harvey D., Heiman T.J., Hernandez J.R., Houck J.,
RA Hostin D., Houston K.A., Howland T.J., Wei M.-H., Ibegwam C.,
RA Jalai M., Kalush F., Karpen G.H., Ke Z., Kennison J.A., Ketchum K.A.
RA Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z.,
RA Lasko P., Lei Y., Levitsky A.A., Li J., Li Z., Liang Y., Lin X.,
RA Liu X., Mattei B., McIntosh T.C., McLeod M.P., McPherson D.,
RA Merkulov G., Milshina N.V., Mobarry C., Morris J., McPherson D.L.,
RA Mount S.M., Moly M., Murphy B., Murphy L., Muzny D.M., Nelson D.L.,
RA Nelson D.R., Nelson K., Nixon K., Nusskern D.R., Pacleb J.M.,
RA Palazzolo M., Pittman G.S., Pan S., Pollard J., Puri V., Reese M.G.,
RA Reinert K., Remington K., Saunders R.D.C., Scheeler F., Shen H.,
RA Spier E., Sjden-Kiamos I., Simpson M., Skupski M.P., Smith T.,
RA Spier E., Spradling A.C., Stapleton M., Strong R., Sun E.,
RA Syirskas R., Tector C., Turner R., Venter E., Wang A.H., Wang Z.-Y., Wassarman D.A., Weinstock G.M., Weissenbach J.,
RA Wang Z.-Y., Wassarman D.A., Weinstock G.M., Weissenbach J.,
RA Yel J., Yeh R.-F., Zaveri J.S., Zhan M., Zhang G., Zhao Q., Zheng I.,
RA Shus B.A., Myers E.W., Rubin G.M., Venter J.C.;
"The genome sequence of Drosophila melanogaster.",
SR EMBL, AB003510, Angarago 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Drosophila melanogaster (Fruit fly).
Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;
Everygota; Meoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
NCBI TaxID=7227;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A. STRAIN-BERKELBY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Q9VWQ3, PRELIMINARY, PRT; 474 AA.
Q9VWQ3, 01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
CG6961 protein.
CG6961.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRINTS; PRUIDUC,
SMART; SM00360; RRM;
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SEQUENCE 474 AA;
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InterPro; IPR000504; RNA_rec_mot.
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V., Reese M.G.,
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                                            RP SEQUENCE FROM N. A.

RC STRAIN-BERKELEY;

RX MEDLINE-20156006; pubMed=10731132;

RA Adams M.D., Celniker S.E., Holt R.A., Evans C.A., Gocayne J.D.,

RA Adams M.D., Celniker S.E., Li P.W., Hoskins R.A., Galle R.F.,

RA Amanatides P.G., Scherer S.E., Li P.W., Hoskins R.A., Galle R.F.,

RA George R.A., Levis S.E., Richards S., Ashburner M., Henderson S.N.,

RA George R.A., Levis S.E., Richards S., Ashburner M., Henderson S.N.,

RA Burandon R.C., Rogers Y.-H.C., Blazej R.G., Champe M., Pfeiffer B.D.,

RA Barandon R.C., Rogers Y.-H.C., Blazej R.G., Champe M., Pfeiffer B.D.,

RA Barandon R.C., Rogers Y.-H.C., Blazej R.G., Champe M., Pfeiffer B.D.,

RA Abril J.F., Agbayani A., An H.-J., Andrews-Pfannkoch C., Baldwin D.,

RA Ballew R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley E.M.,

RA Beesson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S.,

RA Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P.,

RA Burtis K.C., Busam D.A., Butler H., Cadleu E., Center A., Chandra I.,

RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,

RA de Pablos B., Delcher A., Deng Z., Mays A.D., Dew I., Dietz S.M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT
Q9VA50
ID Q9
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Best Local S
Matches 6
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Pterygota; Neopte
Ephydroidea; Dros
NCBI TaxID=7227;
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Q9VA50;
01-MAY-2000
01-MAY-2000
01-JUN-2002
CG9713 prote:
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SMART; SM00360; RRM;
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Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
a; Drosophilidae; Drosophila.
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75.0%;
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Last sequence update)
Last annotation update)
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Pred. No.
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19

Matches Query Match Best Local

RESULT
OPSWOOD
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Q930K0
ID Q930K
AC Q930K
AC Q930K
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DT 01-DE
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DT 01-JU
DT Putat
GN RA019
OG Plasm
OC Bacte
OC Rhizo
OC Bacte
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RA Durbin K.J., Evangelista C.C., Ferrac C., Ferriera S., Fleischmann W.,
RA Fosler C., Gabrielian A.E., Garg N.S., Gelbart W.M., Glasser K.,
RA Glodek A., Gong F., Gorrell J.H., Gu Z., Guan P., Harris M.,
RA Harris N.L., Harvey D., Heiman T.J., Hernandez J.R., Houck J.,
RA Hostin D., Houston K.A., Howland T.J., Wei M.-H., Ibegwam C.,
RA Lin K., Howland T.J., Wei M.-H., Ibegwam C.,
RA Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z.,
RA Liu X., Mattei B., McIntosh T.C., McLeod M.P., McPerson D.,
RA Merkulov G., Milshina N.V., Mobarry C., Morris J., Moshrefi A.,
RA Mount S.M., Molyon K.A., Nixon K., Nusskern D.R., Pacleb J.M.,
RA Melson D.R., Nelson K.A., Nixon K., Nusskern D.R., Pacleb J.M.,
RA Palazzolo M., Pittman G.S., Pan S., Pollard J., Puri V., Reese M.G.,
RA Shue B.C., Siden-Kiamos I., Simpson M., Skupski M.P., Smith T.,
RA Spier E., Spradling A.C., Stapleton M., Skupski M.P., Smith T.,
RA Svirskas R., Tector C., Turner R., Venter E., Wang A.H., Wang X.,
RA Wang Z.-Y., Wassarman D.A., Weinstock G.M., Weissenbach J.,
RA Williams S.M., Woodage T., Worley K.C., Wu D., Yang S., Yao Q.A.,
RA Zheng X.H., Zhong F.N., Zhonn W., Zhang G., Zhao Q., Zheng I.,
RA Zheng R.A., Myers E.W., Rubin G.M., Venter J.C.;
"The genome sequence of Drosophila melanogaster.";
DR RMIL AED01774. AAR7073 1. -
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  Query Match
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EMBL, AE003774, AAF57073.1; --
FlyBase; FBgn0039793; CG9713.
ThrerPro; IPR005123; 20G-FeII Oxy.
ThrerPro; IPR05123; 20G-FeII Oxy.
                                                                                                                                                               MEDLINE=21396509; PubMed=11481432;
Barnett M.J., Fisher R.F., Jones T., Komp C., Abola A.P.,
Barloy-Hubler F., Bowser L., Capela D., Galibert F., Gouzy J.,
Gurjal M., Hong A., Huizar L., Hyman R.W., Kahn D., Kahn M.L.,
Kalman S., Keating D.H., Palm C., Peck M.C., Surzycki R., Wells D.H.,
Yeh K.-C., Davis R.W., Federspiel N.A., Long S.R.;
"Nucleotide sequence and predicted functions of the entire
Sinorhizobium meliloti pSymA megaplasmid.";
Proc. Natl. Acad. Sci. U.S.A. 98:9883-9888 (2001).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RA0195 OR SMA0360.
Rhizobium meliloti (Sinorhizobium meliloti).
Plasmid pSymA (megaplasmid 1).
Bacteria; Proteobacteria; alpha subdivision; Rhizobiaceae group; Rhizobiaceae; Sinorhizobium.
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01-DEC-2001 (TrEMBLrel. 19,
01-JUN-2002 (TrEMBLrel. 21,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCBI_TaxID=382;
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01-DEC-2001
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                                                                                                                Pfam; PF00665; rve; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE
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                                                                                                                                                InterPro;
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                                                        Hypothetical protein;
195 AA; 22083 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FROM N.A.
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                                                                                                                                          IPR001584; Rve.
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85.7%;
  70.6%;
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Pred. No.
  Score 36;
                                                     Complete proteome 4F8481D636BEC9FD
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Length 195;
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X., Smith H.O.,
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                                                                                                                                                                                                                                                                                                                                                01-JUN-2002 (TrEMBLrel. 21, Created)
01-JUN-2002 (TrEMBLrel. 21, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
Hypothetical 50.2 kDa protein.
OSJNBA0042H09.17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Q92XY9;
Q92XY9;
01-DEC-2001
                                                                                                                                                                                                                                           Spermatophyta; Magnoliophyta; Eliliopsida; Ehrhartoideae; Oryzeae; Oryza.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=21396509; PubMed=11481432;
Barnett M.J., Fisher R.F., Jones T., Komp C., Abola A.P.,
Barloy-Hubler F., Bowser L., Capela D., Galibert F., Gouzy J.,
Gurjal M., Hong A., Huizar L., Hyman R.W., Kahn D., Kahn M.L.,
Kalman S., Keating D.H., Palm C., Peck M.C., Surzycki R., Wells D.H.,
Yeh K.-C., Davis R.W., Federspiel N.A., Long S.R.;
"Nucleotide sequence and predicted functions of the entire
Sinorhizobium meliloti pSymA megaplasmid.";
Proc. Natl. Acad. Sci. U.S.A. 98:9883-9888(2001).
EMBL, AB007296; AAK65757.1; -.
InterPro; IPR001647; HTH_TetR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pfam; PF00440; tetR; 1.
Plasmid; Hypothetical protein;
SEQUENCE 201 AA; 21500 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Rhizobium meliloti (Sinorhizobium meliloti).
Plasmid pSymA (megaplasmid 1).
Bacteria; Proteobacteria; alpha subdivision; Rhizobiaceae group; Rhizobiaceae; Sinorhizobium.
                                                           Buell C.R., Yuan Q., Ouyang S., Liu J., Moffat K.S., Hill J.N., Gansberger K., Brenner M., Burgess S., Hance M., Shvartsbeyn M., Tsitrin T., Riggs F., Hsiao J., Zismann V., Blunt S., Pai G., VanAken S.E., Utterback T.R., Feldblyum T.V., Kalb E., Quackenburgsthern S.I. White O. Fracer C.M.
                                                                                                                                                                                                                                                                                                                                           Oryza sativa (Rice).
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                  "Oryza
                    Salzberg S.L., White O., Fraser C.M.; "Oryza sativa chromosome 10 BAC OSJNBa0042H09
                                                                                                                                                                     STRAIN=CV. NIPPONBARE
                                                                                                                                                                                               SEQUENCE
                                                                                                                                                                                                                                         NCBI_TaxID=4530;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-DEC-2001 (TrEMBLrel. 19, 01-JUN-2002 (TrEMBLrel. 21,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TaxID=382;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      46
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  transcriptional regulator.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GHRPLD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GHRPLDKKR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (TrEMBLrel. 19,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Last sequence up
EMBL/GenBank/DDBJ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 36;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
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                                                                                                                                                                                                                                                                                               Embryophyta; Tracheophyta;
a; Poales; Poaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 16; Length 201; 27;
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genomic sequence.";
databases.
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RESULT 14
Q9HA50
ID Q9HA5
AC Q9HA5
DT 01-MA
DT 01-MA
DT 01-MA
DT 01-MA
CDNA
OS HOMO
OC EUKAR
OS HOMO
OC EUKAR
OS HOMO
IS EQUIT
RN [1]
RN [1]
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RN [1]
RA Wagatar
RA Wasar
RA Wasar
RA Watar
RA Wasar
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RA Wanar
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Q8VJW0
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C TISSUB=MAMMARY GLAND;

L ISOJAI T., Ota T., Hayashi K., Sugiyama T., Otsuki T., Suzuki Y.,

A Nishikawa T., Nagai K., Sugano S., Shiratori A., Sudo H.,

A Wagatsuma M., Hosoiri T., Kaku Y., Kodaira H., Kondo H., Sugawara M.,

Takahashi M., Chiba Y., Ishida S., Murakawa K., Ono Y., Takiguchi S.,

A Watanabe S., Kimura K., Murakami K., Ishii S., Kawai Y., Saito K.,

Yamamoto J., Wakamatsu A., Nakamura Y., Nagahari K., Masuho Y.,

A 'Ninomiya K., Iwayanagi T.;

"NEDO human cDNA sequencing project.";

SUbmitted (AUG-2000) to the EMBL/GenBank/DDBJ databases.

BR EMBL, AKO22302; BAB14006.1; -

SO SEQUENCE 505 AA; 54782 MW; 43BADEB48CODFBE7 CRC64;
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Best Local S
Matches 6
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                                                                                               SEQUENCE FROM N.A.

STRAIN-CDC 1551 / OSHKOSH;

Fleischmann R.D., Alland D., Eisen J.A., Carpenter L., White O.,

Fleischmann R.D., Dodson R., Gwinn M.L., Haft D., Hickey E.,

Peterson J., DeBoy R., Dodson R., Ermolaeva M.D., Salzberg S.L.,

Kolonay J.F., Nelson W.C., Umayam L.A., Ermolaeva M.D., Salzberg S.L.,

Delcher A., Utterback T., Weidman J., Khouri H., Gill J., Mikula A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ouery Match 70.6%; Score 36; DB 10; Length 471; Best Local Similarity 100.0%; Pred. No. 63; Matches 6; Conservative 0; Mismatches 0; Indels
laboratory strains.";
Submitted (APR-2001) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-MAR-2002
01-MAR-2002
01-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OBVJWO;
                                                                                                                                                                                                                                                                                                                                 Mycobacterium tuberculosis,
Bacteria; Firmicutes; Actinobacteria; Actinobacteridae;
Actinomycetales; Corynebacterineae; Mycobacteriaceae; Mycobacterium.
                                                           Whole genome comparison of Mycobacterium tuberculosis clinical and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PPE family protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              MT1857
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01-MAR-2001 (TrEMBLrel 16, Last sequence update)
01-MAR-2001 (TrEMBLrel 16, Last annotation update)
CDNA FLJ12240 fis, clone MAMMA1001271.
Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Q9HA50;
01-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=9606;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         41 GHRPLSKQ 48
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity 75.66; Conservative
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2 (TrEMBLrel. 20,
2 (TrEMBLrel. 21,
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Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       694 AA.
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                                                 Query Match
Best Local S
Matches 6
                                                                                   EMBL; AE007044; ARK46130.1; -
TICR; MT1857; -
InterPro; IPR000568; ATPSynt Asub.
InterPro; IPR000030; Microbac_PPE.
Pfam; PF00823; PPE; 1.
PROSITE; PS00449; ATPASE A; UNKNOWN 1.
PROSITE; PS00449; ATPASE A; UNKNOWN 1.
PROSITE; PS00449; ATPASE A; UNKNOWN 1.
85 GHRPCDLRR 93
                      1 GHRPLDKKR 9
                                                 6,
                                                        Similarity
                                               Conservative
                                                      70.6%; Score 36; 66.7%; Pred. No.
                                         Mismatches
                                                       93;
                                                                DB 16;
                                         2; Indels
                                                              Length 694;
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                                     Gaps
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Search comp completed: June me : 23.6 secs 27, 2003, 17:18:05

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Minimum
Maximum
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Maximum Match 100%
Listing first 45 s
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Searched:
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Pred. No.
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                                             A Geneseq 101002:*

1: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1980.DAT:*
2: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1981.DAT:*
3: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1981.DAT:*
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5: /SIDS2/gcgdata/geneseqy-emb1/AA1983.DAT:*
6: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1985.DAT:*
7: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1985.DAT:*
9: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1987.DAT:*
9: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1987.DAT:*
10: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1989.DAT:*
11: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1990.DAT:*
12: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1991.DAT:*
13: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1993.DAT:*
14: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1993.DAT:*
15: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1993.DAT:*
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18: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1993.DAT:*
19: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1993.DAT:*
20: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1993.DAT:*
21: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1993.DAT:*
22: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1993.DAT:*
23: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1993.DAT:*
24: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1993.DAT:*
25: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1993.DAT:*
26: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1993.DAT:*
27: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1993.DAT:*
28: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1993.DAT:*
29: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1993.DAT:*
20: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1993.DAT:*
21: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1993.DAT:*
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51
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Copyright (c) 1993 - 2003 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have score greater than or equal to the score of the result being pland is derived by analysis of the total score distribution. printed,

SUMMARIES

10	9	80	7	6	տ	4	ω	N		Result No.
5 1	51	51	51	51	51	51	51	51	51	Score
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118	87	30	29	28	16	12	12	10	9	Query Match Length DB
21	22	14	14	14	14	14	13	16	22	
AAG00151	ABB11339	AAR42546	AAR42547	AAR40010	AAR44830	AAR44829	AAR28629	AAW11931	ABB76979	ID
Human secreted pro	Human beta-fibrino	Leukocyte-binding	Leukocyte-binding	Scintigraph imagin	Human fibrin beta-	Human fibrin beta-	N-terminal human f	Fibrinogen A-alpha	Residues 45-54 of	Description

New citrulline-containing polypeptide from fibrin, useful for diagnosis and treatment of rheumatoid polyarthritis

WPI; 2001-114394/13.

Serre G,

Sebbag M;

(UYTO-)

UNIV TOULOUSE SABATIER PAUL.

01-JUL-1999; 01-JUL-1999;

99FR-0008470. 99FR-0008470

Example 1; Page 12; 23pp; French.

Homo sapiens	Residue Anti-ar human;	22-JUL-2002	ABB76979	ESULT 1 BB76979 D ABB769		44	43	42	40	9 0	נט נג 7 מ	36	υ <u>υ</u> 5 4	. u	32	3 G 0	29	28	26	25	24	22	21	20	1 8	17	16		14		11	
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	lues 45-54 of arthritic; and fibrinogen	(firs		standard;		70.6	•							•		•			•			80.4	•			100.0	0	0	0 :	30	.0	
1	f human anti-inf	t ent		peptide;		1251	g	505	306	165	153	11	923 11	474	474	5 3 4 9	12	α ο α	ာထ	œ ·	7 ~	7	7	536	4 4 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2	495	491	491	453	141	140	
	fibr flamm chain	ry)		de;		22	22	22																2 1) N	22	22	16	2 1	2 2	22	
•	inogen beta chain atory; fibrin; rhe			9 AA.	ALIGNMENTS	AAB65626	0844	AAB95070	AAY51614	2717	AAB28383	AAB10629	AAB10666	ABB66823	ABB62780	AAU91250	AAR44838	AAR65793	AAR44837	AAP82686	AAR63269	AAR05558	AAP71314	AAM78491	AAM79476	AAM79475	AAM78492	224	AAM78493	л √	ıN	
	in precursor. rheumatoid polyarthritis;					Novel protein kina	human di	Human protein sequ		eptococcus	Sequence block SBW	Fibronectin beta c	prosopnita metanog Fibrinogen-beta pe	Drosophila melanog	Prosophila melanog	Mouse homologue of	Human fibrin beta-	synthetic beta-pep Fibrin-specific ep	Human fibrin beta-	Human fibrin beta	Beta-peptide used Fibrin-specific en	tid	a	Human protein SEO			protein SEQ	_	otein	n polypeptide	Novel human secret	

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RESULT 2
AAW11931
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Best Local S
Matches 9
                                                 Query Match
                                                                                                                                      soluble fibrin in plasma samples (e.g. by sandwich immunoassay) particularly for the diagnosis of pathological conditions such disseminated intravascular coagulation.
                                                                                                                                                                                                            A novel monoclonal antibody (MAb) reacts with human soluble fibrin, but not with human fibrinogen. Specifically when the MAb reacts with urea-treated des-AABB fibrin the reaction is not inhibited by peptides corresponding to fibrinogen A-alpha-chain residues 17-26 (AAWI1931), B-beta-chain residues 15-24 (AAWI1932) or gamma-chain residues 312-324 (AAWI1933). The MAb is useful in immunoassays for collaboration of the collaborat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to a citrulline (Cit) containing polypeptide derived from all or part of the alpha- or beta-chains of fibrin by substitution of at least one arginine residue by Cit. The Cit containing polypeptides can be used for in vitro diagnosis of rheumatoid polyarthritis (RP), by detecting disease-specific autoantibodies, and therapeutically for neutralising the RP-associated autoimmune response. The present sequence is a fragment (residues 45-54) of the human fibringen beta chain precursor, which was used in an example from the
                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Monoclonal antibody with fibrinogen, is
                                                                                                                                                                                                                                                                                                                                                                                    Example 3; Page 15; 32pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                      specimens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1995-206667/27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-NOV-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Monoclonal; antibody; human;
urea-treated; des-AABB; A-al
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Inuzuka K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (IATR ) IATRON LAB INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Fibrinogen A-alpha-chain residues 17-26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     02-APR-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                invention
                        Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                            fibrinogen,
  69; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GHRPLDKKR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ΑA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       intravascular coagulation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first
                                                                                                 A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 93JP-0297325
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             peptide; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                       reactive with soluble human fibrin - but useful for fibrin immunoassay in plasma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kohno
100.0%; S
100.0%; S
tive 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    soluble; fibrin; fibrinogen;
                        Score 51; DB 16;
Pred. No. 0.00058;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 51;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Soe
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           immunoassay; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   7.8e+05
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                                            Length 10;
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Matches

Conservative

Mismatches

Indels

0

Gaps

0

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RESULT 4
AAR44829
ID AAR4
XX
AC AAR4
XX
DT 20-J
XX
DE Huma
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AAR28629
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                                                                                                                                                                                Matches
                                                                                                                                                                                         Query Match
Best Local S
                                                                                                                                                                                                                                             This sequence represents an N-terminal peptide of human fibrin. It was used in the production of bispecific monoclonal antibodies which are specific for fibrins, but do not bind fibrinogen, and and are specific for anti-thrombolytic substance. The compsn. contg. these Abs lacks the side effects of prior art Ab targetted thrombolytic agents and has enhanced thrombolytic activity.
           Human fibrin
                                                                                                                                                                                                                                                                                                                                                            Bispecific antibody useful for treating thrombotic obstructive diseases e.g. cardiac infarction - comprises antithrombus antibody variable region and anti-thrombolyric substance antibody variable region with no heavy chain constant region domains 2 and
                                  20-JUN-1994
                                                                             AAR44829
                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                            Disclosure; Page 3; 30pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1992-383677/47
                                                                                                                                                                                                                                                                                                                                                                                                                                           Iwasa S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17-MAY-1991;
13-MAR-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14-MAY-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EP513778-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               peripheral arterial/venous obstruction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        bispecific hybrid monoclonal antibody; thrombolytic agent; cardiac infarction; arterial embolism; cerebral infarction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           N-terminal human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAR28629;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (TAKE ) TAKEDA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19-NOV-1992.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22-MAR-1993
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                                                                                                                                                        1 GHRPLDKKR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    μ
                                                                                                                                                                               Similarity 9; Conserv
                                                                            standard; peptide; 12
                                                                                                                                    GHR PLDKKR
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Kurokawa T,
                                                                                                                                                                                                                           12 AA;
                                                                                                                                                                                Conservative
           beta-chain N-terminal
                                (first entry)
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92JP-0055025
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         fibrin peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    9
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Watanabe
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                                                                                                                                                                                         Score
Pred.
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          peptide (1-11)-Cys
                                                                                                                                                                                         No.
                                                                                                                                                                                          0.00071;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               retinal
                                                                                                                                                                                                     BB
                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               arterial obstruction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         infarction;
                                                                                                                                                                                                    Length
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RESULT 5
AAR44830
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                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tissue plasminogen activator; t-PA; mutein; fibrin; antigen; anti-fibrin; monoclonal antibody; hybridoma; thrombolysis antithrombotic agent; bispecific antibody.
                                                                                                                                 Tissue plasminogen activator; t-PA; mutein; fibrin; antigen; anti-fibrin; monoclonal antibody; hybridoma; thrombolysis antithrombotic agent; bispecific antibody.
                                                                                                                                                                                Human fibrin
                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                    Human fibrin beta-chain N-terminal peptide (1-11)-Cys was synthesised and coupled to BSA for injection into mice. The peptide was used to raise antibodies to human fibrin. Monoclonal antibodies specific for fibrin are used in the production of bispecific monoclonal antibodies which also recognise truncated tPA muteins lacking the finger, EGF and Kringle 1 domains.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 9; 38pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hybrid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17-JUN-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                       JP05304992-A
                                                                                                                                                                                                                                                 AAR44830 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20-JUN-1991;
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                                                                                                             Synthetic
                                                                                                                                                                                                      20-JUN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         reduced reactivity to fibrinogen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1993-408334/51.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (TAKE ) TAKEDA CHEM IND
(TAKE ) TAKEDA CHEM IND LTD
                      20-JUN-1991;
                                            17-JUN-1992;
                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    monoclonal antibody - used for prepn. of thrombolytic drug increased thrombolytic activity and specificity and
                                                                                                                                                                                                                                                                                                                             1 GHRPLDKKR
                                                                                                                                                                                                                                                                                                                                                                                                12
                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                              beta-chain internal peptide fragment.
                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 91JP-0148936
                      91JP-0148936
                                            92JP-0158301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        92JP-0158301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note=
12
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                                                                                                                                                                                                                                                                                                                              9
                                                                                                                                                                                                                                                 peptide; 16
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                                                                                                                                                                                                                                                                                                                                                               100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "BSA carrier is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "human fibrin beta-chain
                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                                                                                                                                  Score 51; DB 14;
Pred. No. 0.00071;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                residues 1-11"
                                                                                                                                                                                                                                                                                                                                                                         Length 12
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RESULT 6
AAR40010
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Query Match
Best Local Similarity
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Best Local
                                               The sequence is that of a specific binding peptide used as part of a reagent for preparing a scintigraphic imaging agent for imaging sites within a mammalian body. In this the peptide is covalently linked to a radiolabel-binding moiety which is capable of forming a complex with a radioisotope, pref. technetium-99M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human fibrin internal peptide fragment was identified as a candidate immunogen to raise antibodies to human fibrin. Monoclonal antibodies specific for fibrin are used in the production of bispecific monoclonal antibodies which also recognise truncated tPA muteins
                                                                                                                                    Reagents for preparing scintigraphic imaging agents - technetium-99M labelled peptide(s) contg. 3-100 aminos
                              Sequence
                                                                                                                 Claim 35;
                                                                                                                                                                    WPI; 1993-368429/46
                                                                                                                                                                                        Buttram S,
                                                                                                                                                                                                                                 30-APR-1992;
                                                                                                                                                                                                                                                                           11-NOV-1993.
                                                                                                                                                                                                                                                                                                WO9321962-A.
                                                                                                                                                                                                                                                                                                                              Modified-site
                                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                Reagent; site imaging; technetium-99M labelled; peptide
                                                                                                                                                                                                                                                                                                                                                                                                    Scintigraph imaging agent specific binding peptide
                                                                                                                                                                                                                                                                                                                                                                                                                          23-MAY-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                              AAR40010
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 9; 38pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               reduced
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                                                                                                                                                                                                            (DIAT-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            increased thrombolytic activity reactivity to fibrinogen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        monoclonal antibody - used for prepn. of thrombolytic drug increased thrombolytic activity and specificity and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity 9; Conserv
                                                                                                                                                                                                            DIATECH INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GHRPLDKKR
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                                                                                                               Page 39; 55pp; English.
                               28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EGF and Kringle 1 domains.
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100.0%;
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Pred.
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Pred.
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No.
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DB 14;
0.0018;
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aminoacid(s)
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Matches

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AAR42546 standard; Peptide; 30
                                                                                                                                                                                                                             New peptides are claimed which are leukocyte binding peptides having covalently bound to them a molety which can bind a Tc-99m radiolabel. The peptides having Tc-99m bound to them are useful & scintigraphic imaging agents for imaging sites of infection and inflammation in the mammalian body, e.g. caused by ischaemia, inflammatory bowel disorder, arthritis or tumours. The present sequence is a specifically claimed example of such a peptide derived from fibrinopeptide B chain.
                                                                                                                                                                                                                                                                                                                                                                   Claim 15; Page 31; 40pp;
                                                                                                                                                                                                                                                                                                                                                                                                   Scintigraphic imaging agent for sites of inflammation -
comprising leukocyte-binding peptide bound technetium-99m via
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Leukocyte binding peptide; fibrinopeptide B chain; scintigraphic imaging; inflammatiom site; technetium 99m.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Leukocyte-binding peptide which can bind to technetium-99m
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAR42547 standard; Peptide; 29 AA.
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                                                                                                                                                          100.0%; Score 51; DB 14; llarity 100.0%; Pred. No. 0.0019; Conservative 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                93WO-US02320
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /label= OTHER
/note= "BAT, i.e. N6,N9-bis(2-methyl-
2-mercaptopropyl)-6,9-diazanonanoic acid"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note= "Arg-NH2"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                   English
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                                                                                                                                                        Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                              Length 29;
                                                                                                                                                    Indels
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                                                                                                                                               Gaps
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Best Local S
Matches 9
Human; cytokine; cell proliferation; cell differentiation; growth factor; haematopoiesis regulation; tissue growth; immunomodulator; activin; inhibin; chemotaxis; chemokinesis; thrombolysis; oncogenesis; proliferation; metastasis; cancer; tumour; haematopoietic disorder; myeloid cell disorder; lymphoid cell disorder; asthma; arthritis;
                                                                                        Human beta-fibrinogen homologue,
                                                                                                                            11-JAN-2002
                                                                                                                                                                                 ABB11339
                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                            New peptides are claimed which are leukocyte binding peptides having covalently bound to them a moiety which can bind a Tc-99m radiolabel. The peptides having Tc-99m bound to them are useful scintigraphic imaging agents for imaging sites of infection and inflammation in the mammalian body. e.g. caused by ischaemia, inflammatory bowel disorder, arthritis or tumours. The present sequence is a specifically claimed example of such a peptide derived from fibrinopeptide B chain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Scintigraphic imaging agent for sites of inflammation -
comprising leukocyte-binding peptide bound technetium-99m via
binding moiety
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 15; Page 31; 40pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1993-303154/38.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Buttram S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (DIAT-) DIATECH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13-MAR-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12-MAR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16-SEP-1993.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO9317719-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Leukocyte binding peptide; fibrinopeptide B chain; scintigraphic imaging; inflammatiom site; technetium 99m.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Leukocyte-binding peptide which can bind to technetium-99m
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05'-AUG-1994
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                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                standard;
                                                                                                                                                                                                                                                                             GHRPLDKKR 9
                                                                                                                                                                                                                                                   GHRPLDKKR
                                                                                                                                                                                                                                                                                                                                                                30 AA;
                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                              peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note= "acetamidomethyl-Cys"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         note=
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                                                                                                                                                                                                                                                                                                                    100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Lees RS,
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                                                                                                                                                                              87
                                                                                                                                                                                                                                                                                                                 Score 51;
Pred. No.
                                                                                                                                                                             A
                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                         SEQ ID NO:1709
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Lister-James
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W09317719-A

Modified-site

Synthetic

05-AUG-1994

AAR42547;

Modified-site

WPI; 1993-303154/38.

moiety

Buttram S, (DIAT-) 13-MAR-1992; 12-MAR-1993;

Dean RT,

DIATECH INC.

RESULT 8
AAR42546
ID AAR4

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GHRPLDKKR

10

GHRPLDKKR 9

Matches Query Match

Local

Similarity 9; Conser

Sequence

Query Match

22;

Length

87;

chronic inflammatory condition; proliferative retinopathy; atherosclerosis; coronary heart disease; arterial ischaemia; bone disorder; osteoporosis; vascular growth disorder; tissue regeneration; wound healing; infection; immune disorder; cell culture; drug screening; gene therapy; antiinflammatory; antiasthmatic; antiarthritic; haemostatic; antiarteriosclerotic; cytostatic; osteopathic; vasotropic; cardiant; virucide; antibac antifungal; vulnerary; antiulcer. antibacterial;

Homo sapiens.

WO200157188-A2

05-FEB-2001; 2001WO-US03800

03-FEB-2000; 27-APR-2000; 2000US-0496914. 2000US-0560875.

(HYSE-) HYSEQ INC

Liu C, Drmanac RT

WPI; 2001-457740/49

Human proteins and DNA encoding sequences useful for preventing, treating or ameliorating a medical condition in a mammalian subject e.g. arthritis and cancer - $\,$

Claim 20; Page 167; 1963pp; English.

sequences ABA08225-ABA09574 represent nucleic acids encoding them. The cinvention also relates to vectors and recombinant host cells comprising a composition also relates to vectors and recombinant host cells comprising a contribution also relates to vectors and recombinant host cells comprising a contribution also relates the polypeptides, methods of detecting the nucleotides or polypeptides in a sample, and methods of detecting the nucleotides or polypeptides of the invention have homology to known proteins, thereby compounds which compounds an insight into their probable biological activities, and hence polypeptides of the invention. Although novel, many of the polypeptides of the invention have homology to known proteins, thereby conditions activities, including cytokine, cell proliferation or cell differential therapeutic applications. The polypeptides of the invention may have various activities, including cytokine, cell proliferation or cell chamanomodulatory activity; activin- or inhibin-related activities; chemotactic or chemokinetic activities; haemostatic, thrombotic or thrombotytic activities; activin- or inhibin-related activities; chemotactic or chemokinetic activities; haemostatic, thrombotic or thrombotytic activities; receptor or ilgand activities; or may be convolved in oncogenesis, cancer cell proliferation or metastasis.

CC pepending on their biological activities, polypeptides and nucleotides of the invention are useful for preventing, treating or ameliorating medical conditions, e.g., by protein or gene therapy. Such conditions include cancers, haematopoietic disorders (e.g., myeloid or lymphoid cell conditions, e.g., by protein or gene therapy. Such conditions and cell conditions and abnormal cell is activities and biological activities, and abnormal cell called the provides and conditions and ulcers), while those with cell cell infections in addition to immune disorders. Polypeptides with growth factor activity may be used in cell cultures to promote cell growth. For example, such polypeptides may be used to manipulate stem cells in culture to give rise to neuroepithelial cells that can be used to augment or replace cells damaged by illness, autoimmune disease or accidental damage. The polypeptides and nucleotides may also be used in the diagnosis of the above conditions, and in drug screening techniques. The present sequence represents a novel human Sequences ABB10981-ABB12330 techniques. de of the in represent 1350 novel human polypeptides, represent nucleic acids encoding them. The

RESULT 11
AAU33273
ID AAU3:

AAU33273 standard; Protein;

140 AA

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45

GHRPLDKKR

53

1 GHRPLDKKR

Matches

Similarity 9; Conserv

Conservative

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0.0088;

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RESULT 10
AAG00151
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Query Match
Best Local S
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                                                                                                         The present sequence is a polypeptide encoded by one of a large number of 5' ESTs derived from mRNAs encoding secreted proteins. The 5' ESTs were prepared from total human RNAs or polyA+ RNAs derived from 30 different tissues. EST sequences usually correspond mainly to the 3' untranslated region (UTR) of the mRNA because they are often obtained from oligo-dT primed cDNA libraries. Such ESTs are not well suited for isolating cDNA sequences derived from the 5' ends of mRNAs and even in those cases where longer cDNA sequences have been obtained, the full 5' UTR is rarely included. 5' ESTs are derived from mRNAs with intact. 5' ends and can therefore be used to obtain full length cDNAs and genomic
                                                                   DNAs. 5' ESTs are also used in diagnostic, forensic, gene therapy and chromosome mapping procedures. They are used to obtain upstream regulatory sequences and to design expression and secretion vectors.
                                          Sequence
                                                                                                                                                                                                                                                                                                        New nucleic acid that is a 5' expressed sequence tag (5' EST) for obtaining cDNAs and genomic DNAs that correspond to 5'ESTs and fo
                                                                                                                                                                                                                                                                Claim 13; SEQ ID 4232; 71pp + CD-ROM; English.
                                                                                                                                                                                                                                                                                           diagnostic, forensic, gene
                                                                                                                                                                                                                                                                                                                                                   N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                WPI; 2000-500381/45.
                                                                                                                                                                                                                                                                                                                                                                                           Dumas Milne Edwards
                                                                                                                                                                                                                                                                                                                                                                                                                                              26-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21-FEB-2000; 2000EP-0200610
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-SEP-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
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                                         118 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      chromosome
                                                                                                                                                                                                                                                                                                                                                                                                                                                99US-0122487
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  expressed sequence tag; secreted protein; cDNA isolation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               protein,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     mapping.
                                                                                                                                                                                                                                                                                           therapy and
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  Score
Pred.
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Pred. No. 0.0063
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NO: 4232.
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  No.
                                                                                                                                                                                                                                                                                                                                                                                         Giordano
                                                                                                                                                                                                                                                                                           chromosome mapping procedures
                 DΒ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
              Length 118;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
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RESULT 12
AAO10778
ID AAO10
XX AAO10
AC AAO10
DT 06-NO
XX Human
XX Human
                                                                                                                                                                                                                                                                                                 The invention relates to novel human secreted polypeptides. The CC polypeptides and antibodies to the polypeptides are useful for CC determining the presence of or predisposition to a disease associated (CC with altered levels of polypeptide. The polypeptides are also useful for identifying agents (agonists and antagonists) that bind to them. Cells for use in treatment of a pathology related to aberrant expression or CC the nucleic acids encoding the polypeptides. Vectors comprising CC The proteins are useful for polypeptides and cells genetically CC therapeut to express them are also useful for producing the proteins. CC increase stem cell in genetic vaccination, testing and consequence of the proteins are useful in genetic vaccination, testing and CC increase stem cell proliferation; to regulate haematopoiesis; and in CC in treatment of leukaemias. AMU29510-AMU3304 represent the amino acid consequences of novel human secreted proteins of the invention.
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                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                            Query Match
Human; cytokine; cell proliferation; cell differentiation; gene therapy;
                                        Human polypeptide SEQ ID NO 24670.
                                                                            06-NOV-2001
                                                                                                      AA010778;
                                                                                                                    AAO10778 standard; Protein; 141 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 20; Page 752; 765pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nucleic acids encoding a range of human polypeptides, useful in genetic vaccination, testing and therapy -
                                                                                                                                                                                                                                                Local Similarity
les 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-611725/70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tang YT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-APR-2000; 2000US-0552929.
26-JAN-2001; 2001US-0770160.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16-APR-2001; 2001WO-US08656.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-OCT-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200179449-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; vaccination; gene therapy; nutritional supplement; stem cell proliferation; haematopoiesis; nerve tissue regeneration; immune suppression; immune stimulation; anti-inflammatory; leukaemi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel human secreted protein #3764
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAU33273;
                                                                                                                                                                                            45 GHRPLDKKR
                                                                                                                                                                                                                     GHRPLDKKR 9
                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Liu C, Drmanac
                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                         100.0%;
                                                                                                                                                                                                                                            ,
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                                                                                                                                                                                                                                               Score 51; DB 22;
Pred. No. 0.011;
                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                      0,
                                                                                                                                                                                                                                                          Length 140;
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RESULT 13
AAG00150
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Best Local Similarity
Yarches 9; Conserv;
                                                                                                                                                                                                                                                                                                                                              EP1033401-A2
                        Homo sapiens
                                                                              Human secreted protein, SEQ ID NO: 4231.
                                                      gene
                                                                Human; 5' EST;
                                                                                                                    06-OCT-2000
                                                                                                                                              AAG00150;
                                                                                                                                                             AAG00150 standard; Protein; 150 AA
                                                                                                                                                                                                                                                                                                                                                                                             production of other cytokines in other cell populations. The polynucleotides and polypeptides are useful in gene therapy, vaccines or e.g. stem cell growth factor activity, haematopoiesis regulating activity tissue growth factor activity, haematopoiesis regulating activity infinibin activity and may be useful in the diagnosis and/or treatment of cancer, leukaemia, nervous system disorders, arthritis and
                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                  Note: The sequence data for this patent did not specification, but was obtained in electronic fo at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to human polynucleotides (AAI79941-AAI93841) and the encoded proteins (AAO000010-AAO13910) that exhibit activity elating to cytokine, cell proliferation or cell differentiation or which may induce
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 20; SEQ ID NO 24670; 1399pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Isolated nucleic acids and polypeptides, useful for preventing diagnosing and treating e.g. leukaemia, inflammation and immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-514838/56.
N-PSDB; AAI90709.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               vaccine; peptide therapy; stem cell growth factor; haematopoiesis;
tissue growth factor; immunomodulatory; cancer; leukaemia;
nervous system disorders; arthritis; inflammation.
                                                  therapy;
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                                                                                                                                                                                                                                  41 GHRPLDKKR 49
                                                                                                                                                                                                                                                     1 GHRPLDKKR
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                                                                                                                                                                                                                                                                                                                                     141 AA;
                                                                                                                                                                                                                                                                                   100.0%;
ilarity 100.0%;
Conservative 0
                                                                                                           (first entry)
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                                   expressed sequence tag; secreted protein; chromosome mapping.
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Pred. No. 0.011;
                                                                                                                                                                                                                                                                             Mismatches
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                                                                                                                                                                                                                                                                                                Length 141;
                                                                                                                                                                                                                                                                         Indels
                                            cDNA isolation,
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RESULT 14
AAM78493
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Best Local
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3-FEB-2000;
27-APR-2000;
20-JUN-2000;
19-JUL-2000;
01-SEP-2000;
15-SEP-2000;
20-OCT-2000;
30-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    of 5' ESTs derived from mRNAs encoding secreted proteins. The 5' ESTs were prepared from total human RNAs or polyA+ RNAs derived from 30 different tissues. EST sequences usually correspond mainly to the 3' untranslated region (UTR) of the mRNA because they are often obtained from oligo-dT primed cDNA libraries. Such ESTs are not well suited for isolating cDNA sequences derived from the 5' ends of mRNAs and even in those cases where longer cDNA sequences have been obtained, the full 5' UTR is rarely included. 5' ESTs are derived from mRNAs with intact 5' ends and can therefore be used to obtain full length cDNAs and genomic DNAs. 5' ESTs are also used in diagnostic, forensic, gene therapy and chromosome mapping procedures. They are used to obtain upstream
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New nucleic acid that is a 5' expressed sequence tag (5' EST) obtaining cDNAs and genomic DNAs that correspond to 5'ESTs and diagnostic, forensic, gene therapy and chromosome mapping pro-
                                                                                                                               05-FEB-2001;
                                                                                                                                                            09-AUG-2001.
                                                                                                                                                                                                                                            Human; cytokine; cell proliferation; cell differentiation; gene th vaccine; peptide therapy; stem cell growth factor; haematopoiesis; tissue growth factor; immunomodulatory; cancer; leukaemia; nervous system disorder; arthritis; inflammation.
                                                                                                                                                                                                                                                                                                                                                   06-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                          AAM78493 standard; Protein; 453
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present sequence is a polypeptide encoded by one of a large of 5' ESTs derived from mRNAs encoding secreted proteins. The 5'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 13; SEQ ID 4231; 71pp + CD-ROM; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        N-PSDB;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-FEB-1999;
                                                                                                                                                                                         WO200157190-A2.
                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                    Human protein SEQ ID NO 1155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21-FEB-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              regulatory sequences and to design expression and secretion vectors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   45
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GHRPLDKKR 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GHRPLDKKR 53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 150 AA;
                                                       2000US-0496914.
2000US-0560875.
2000US-0598075.
2000US-0620325.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
 2000US-0663561.
2000US-0693325.
2000US-0728422.
                                                                                                                               2001WO-US04098
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                                                                                                                                                                                                                                                                                                                                                (first entry)
                                             2000US-0654936.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            99US-0122487
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%;
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Pred. No. 0.011;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        mapping procedures
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RESULT 15
AAR82243
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local
                                                                                                                                                                                                                             Human fibrinogen B-beta chain; transgenic animal milk; treatment; sheep beta-lactoglobulin signal peptide fusion protein; surgical adhesive.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     polynucleotides and polypeptides are useful in gene therapy, vaccines or peptide therapy. The polypeptides have various cytokine-like activities, e.g. stem cell growth factor activity, haematopoiesis regulating activity, tissue growth factor activity, immunomodulatory activity and activity/inhibin activity and may be useful in the diagnosis and/or treatment of cancer, leukaemia, nervous system disorders, arthritis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Tang YT,
Zhao QA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to polynucleotides (AAK51456-AAK53435) and the encoded polypeptides (AAM78333-AAM80302) that exhibit activity elating to cytokine, cell proliferation or cell differentiation or which may induce production of other cytokines in other cell populations. The
          N-PSDB; AAT03852
                      WPI; 1995-320582/41
                                             Dalrymple MA,
                                                                                                         03-MAR-1994;
                                                                                                                                01-MAR-1995;
                                                                                                                                                         08-SEP-1995
                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                Human
                                                                                                                                                                                                                                                                                                                                                        AAR82243 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Note: Records for SEQ ID NO 2110 (AAK52581), (AAM80020) are omitted as the relevant pages
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 20; Page 3390-3391; 6221pp; English
                                                                     (PHAR-)
                                                                                                                                                                                WO9523868-A1
                                                                                                                                                                                                                                                                                                        26-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         were missing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucleic acids encoding polypeptides with cytokine-like activities, useful in diagnosis and gene therapy -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-476283/51.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              inflammation.
                                                                                                                                                                                                                                                                             fibrinogen B-beta chain protein
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5
                                                                                                                                                                                                                                                                                                                                                                                                                                           1 GHRPLDKKR
                                                                    PHARM PROTEINS LTD ZYMOGENETICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HYSEQ INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                   GHRPLDKKR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Liu C, Drmanac RT, Asundi V,
Wang D, Wang J, Zhang J, Ren
Yang Y, Wejhrman T, Goodrich R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  453
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          at
                                                                                                                                                                                                                                                                                                      (first entry)
                                               Foster DC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AA;
                                                                                                         94US-0206176
                                                                                                                                 95WO-US02648
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          time
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100.0%; Pred. No. 0.039;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         of publication
                                             Garner I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
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                                               Prunkard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 22;
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F, C
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from
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 453;
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the sequence listing
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В δ

DNA

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RESULT 16
AAM/78492
AID AAM/78-
XX AAM/78-
XX AAM/78-
XX AAM/78-
XX AAM/78-
XX Human
XW Human
XW Human
XW Homo
XX Homo
XX PO-AU
XX O5-FE
PR 05-FE
PR 27-AP
PR 19-JU
PR 119-JU
PR 119-JU
PR 119-JU
PR 119-JU
PR 12-AP
PR 20-OC
XX O1-SE
PR 19-OC
XX O1-SE
PR 20-OC
XX O1-SE
PR 20-OC
XX O1-SE
PR 10-SE
PR 10-
                                                                                                                                                                                                                                                                                                 03-FEB-2000; 2000US-0496914.

27-APR-2000; 2000US-0560875.

20-JUN-2000; 2000US-0598075.

19-JUL-2000; 2000US-0620325.

01-SEP-2000; 2000US-063561.

15-SEP-2000; 2000US-0635325.

20-OCT-2000; 2000US-0693325.
               Claim 20; Page 3389-3390; 6221pp; English
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Best Local
                                                           Nucleic acids encoding puseful in diagnosis and
                                                                                                                                                                                                                                                                                         30-NOV-2000;
                                                                                                                                                                                                                                                   (HYSE-) HYSEQ INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               09-AUG-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; Cytokine; cell proliferation; cell differentiation; gene therapy; vaccine; peptide therapy; stem cell growth factor; haematopoiesis; tissue growth factor; immunomodulatory; cancer; leukaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    06-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAM78492;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAM78492 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              preferably by sheep beta-lactoglobulin signal peptide fusion preferably by sheep beta-lactoglobulin signal peptide fusion preferably of a sheep, pig, goat or cattle. Co-expression with the fibrinogen A-alpha chain (see AAR82244) and the gamma chain (see useful in human and veterinary medicine, e.g. in the formulation of surgical adhesives, which also consist of Factor-XIII, and as a coating surface for polymeric articles, such as synthetic vascular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human fibrinogen B-beta chain may be produced recombinantly, preferably by sheep beta-lactoglobulin signal peptide fusion
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                                                                                                                                     2001-476283/51.
                                                                                                                       AAK51625
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein SEQ ID NO 1154.
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                                                                                                                                                                           Yang
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                                                                                                                                                                 Liu C, Drmanac RT, Asundi V,
Wang D, Wang J, Zhang J, Re
Yang Y, Wejhrman T, Goodrich
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ilarity 100.0%;
Conservative
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                                               polypeptides with cytokine-like
d gene therapy -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             arthritis; inflammation.
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Pred. No. (
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F, (
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Wang
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ng ZW;
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RESULT 17
AAM79475
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N-PSDB;
                                                 Nucleic
useful i
                                                                                                                                                           Tang YT,
Zhao QA,
Xue AJ,
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Best Local
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27-APR-2000; 2000US-0560875.
20-UUN-2000; 2000US-0598075.
19-UUL-2000; 2000US-0620325.
01-SEP-2000; 2000US-063561.
15-SEP-2000; 2000US-063325.
20-OCT-2000; 2000US-0693325.
30-NOV-2000; 2000US-0728422.
Claim 20; Page 257-258; 6221pp; English.
                                                                                                                                                                                               (HYSE-) HYSEQ INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; cytokine; cell proliferation; cell differentiation; vaccine; peptide therapy; stem cell growth factor; haematop tissue growth factor; immunomodulatory; cancer; leukaemia; nervous system disorder; arthritis; inflammation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to polynucleotides (AAK51456-AAK53435) and the encoded polypeptides (AAM78323-AAW80302) that exhibit activity elating to cytokine, cell proliferation or cell differentiation or which may induce production of other cytokines in other cell populations. The polynucleotides and polypeptides are useful in gene therapy, vaccines or e.g. stem cell growth factor activity, haematopoiesis regulating activity, tissue growth factor activity, haematopoiesis regulating activity, inthibin activity and may be useful in the diagnosis and/or activity and may be useful in the diagnosis and/or activity.
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                                                                                     2001-476283/51
)B; AAK52608.
                      c acids encoding polypeptides with cytokine-like activities,
in diagnosis and gene therapy. -
                                                                                                                               Liu C,
Wang D,
Yang Y,
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9; Conserv
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D, Wang J, Zh
, Wejhrman T,
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                                                                                                                         c RT, Asundi V, 2
J, Zhang J, Ren H
an T, Goodrich R;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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F, C
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                                                                                                                                        u P,
Chen
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                                                                                                                       u C,
Wang
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RESULT 18
AAM79476
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Best Local
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27-APR-2000;
20-JUN-2000;
19-JUL-2000;
01-SEP-2000;
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20-OCT-2000;
30-NOV-2000;
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                                                                                                                                                Tang
Zhao
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; cytokine; cell proliferation; cell differentiation; gene th vaccine; peptide therapy; stem cell growth factor; haematopoiesis; tissue growth factor; humnomodulatory; cancer; leukaemia; nervous system disorder; arthritis; inflammation.
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                         Claim
                                                Nucleic acids encoding polypeptides with useful in diagnosis and gene therapy -
                                                                                                                                                                                                                                                                                                                                                                    09-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human protein SEQ ID NO 3122
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                                                                                          WPI; 2001-476283/51
N-PSDB; AAK52609.
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                                                                                                                                             E, G
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                      20; Page 258; 6221pp; English
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                                                                                                                                Yang
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                                                                                                                                Liu C,
Wang D,
Yang Y,
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2000US-0693325.
2000US-0728422.
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2000US-0598075.
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2000US-0654936.
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                                                                                                                                Drmanac RT, Asundi V,
Wang J, Zhang J, Ren
Wejhrman T, Goodrich R,
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Pred. No.
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Wang
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The invention relates to polynucleotides (AAK51456-AAK53435) and the

The invention relates to polynucleotides encoded polypeptides (AAM78323-AAM80302)

(AAK51456-AAK53435) and the that exhibit activity elating

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RESULT 19
AAM79477
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Matches 9
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27-APR-2000; 2000US-0598075.

20-JUN-2000; 2000US-0598075.

19-JUL-2000; 2000US-0620325.

01-SEP-2000; 2000US-0654936.

15-SEP-2000; 2000US-0663561.

20-OCT-2000; 2000US-0693325.

30-NOV-2000; 2000US-0728422.
                                                                                                                              Tang YT,
Zhao QA,
Xue AJ,
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                                                                                                                                                                                                                                                                                                                                                                                                                                Human; cytokine; cell proliferation; cell differentiation; gene therapy; vaccine; peptide therapy; stem cell growth factor; haematopoiesis; tissue growth factor; immunomodulatory; cancer; leukaemia;
                                                                                            N-PSDB;
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                                 Claim 20; Page
                                                                   Nucleic
                                                                                            WPI; 2001-476283/51.
N-PSDB; AAK52610.
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                                                        acids encoding polypeptides with in diagnosis and gene therapy -
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                                 258; 6221pp; English.
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Pred. No.
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                                                                     cytokine-like activities,
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F,
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RESULT 20
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19-JUL-2000;
01-SEP-2000;
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Zhao
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Nucleic acids encoding polypeptides with cytokine-like activities, useful in diagnosis and gene therapy -
                                                                                                                 WPI; 2001-476283/51.
N-PSDB; AAK51624.
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27-APR-2000;
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20-OCT-2000;
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2000US-0620325.
2000US-0654936.
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2000US-0693325
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Wang J, Zhang J, Ren F, (
Wejhrman T, Goodrich R;
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The invention relates to polynucleotides (AAK51456-AAK53435) and the encoded polypeptides (AAM78323-AAM80302) that exhibit activity elating to cytokine, cell proliferation or cell differentiation or which may induce

Claim 20; Page 3388-3389;

6221pp; English.

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Matches
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     inflammation. Note: Records for SEQ ID NO 2110 (AAK52581), 2111 (AAM80020) are omitted as the relevant pages from were missing at the time of publication.
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Search completed: June 29, 2003, 09:59:29 Job time: 72 secs

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Perfect score:
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seq length: 2000000000
     protein search, using sw model
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1: /cgn2_6/ptcdata/1/pubpaa/US08_NEW_PUB.pep:*

2: /cgn2_6/ptcdata/1/pubpaa/US06_NEW_PUB.pep:*

3: /cgn2_6/ptcdata/1/pubpaa/US06_NEW_PUB.pep:*

4: /cgn2_6/ptcdata/1/pubpaa/US07_NEW_PUB.pep:*

5: /cgn2_6/ptcdata/1/pubpaa/US07_NEW_PUB.pep:*

6: /cgn2_6/ptcdata/1/pubpaa/US07_PUBCOMB.pep:*

7: /cgn2_6/ptcdata/1/pubpaa/US07_PUBCOMB.pep:*

9: /cgn2_6/ptcdata/1/pubpaa/US09_PUBCOMB.pep:*

9: /cgn2_6/ptcdata/1/pubpaa/US09_PUBCOMB.pep:*

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US-10-142-935-6
US-10-142-935-4
US-10-131-543-5
US-10-131-346-5
US-10-131-546-5
US-10-131-546-5
US-10-113-724-6
US-10-1142-935-9
US-10-142-935-9
US-10-142-935-5
US-10-123-512-296
US-10-123-512-296
US-10-123-512-296
US-10-133-512-296
US-10-133-512-296
US-10-133-512-296
US-10-133-512-296
US-10-133-512-296
US-10-133-513-34308
US-10-130-8758-11
US-10-130-8758-11
US-10-130-8758-11
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US-10-142-935-6
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ALIGNMENTS

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Query Match
Best Local Similarity
Watches 9; Conserve
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US-10-142-935-4
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Publication No. US20030044418A1
GENERAL INFORMATION:
APPLICANT: DAVIS, Stacey
APPLICANT: HOOK, Magnus A.O.
TITLE OF INVENTION: METHOD AND COMPOSITIONS FOR INHIBITING THROMBIN-INDUCED COAGULAT
FILE REFERENCE: P07201US01/BAS
                                                                                                                                                Sequence 4, Application US/10142935
Publication No. US20030044418A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/10/142,935
CURRENT FILING DATE: 2002-05-13
PRIOR APPLICATION NUMBER: US 60/290,072
PRIOR FILING DATE: 2001-05-13
NUMBER OF SEQ ID NOS: 11
SOFTWARE: PatentIn version 3.1
SEQ ID NO 6
                APPLICANT: DAVIS, Stacey
APPLICANT: HOOK, Magnus A.O.
APPLICANT: HOOK, MAGNUS A.O.
TITLE OF INVENTION: METHOD AND COMPOSITIONS FOR INHIBITING THROMBIN-INDUCED COAGULAT
FILE REFERENCE: PO7201US01/BAS
CURRENT APPLICATION NUMBER: US/10/142,935
CURRENT FILING DATE: 2002-05-13
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TYPE: PRT
PRIOR APPLICATION NUMBER: US 60/290,072
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DB 9; Length 30;

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FILE REFERENCE: 09744-017001

CURRENT APPLICATION NUMBER: US/10/131,346

PRIOR APPLICATION VIMBER: US/05/3500

PRIOR APPLICATION NUMBER: US 09/695,360

PRIOR FILING DATE: 2000-10-24

PRIOR PRIOR PRIOR OF THE CONTROL 
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NAME/KEY: AMIDATION
LOCATION: 30
US-10-131-543-5
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                                                                                                                                                                                                                                                                                                                                                                           Sequence 5, Application US/10131346
Publication No. US20030103899A1
GENERAL INFORMATION:
APPLICATE CONT.
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US-10-131-543-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                        APPLICANT: Cyr, John E.
TITLE OF INVENTION: STABILIZATION OF RADIOPHARMACEUTICAL COMPOSITIONS
TITLE OF INVENTION: USING HYDROPHILIC 6-HYDROXY CHROMANS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 29
SOFTMARE: FastSEQ for Windows Version 4.0
SEQ ID NO 5
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Best Local S
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PRIOR APPLICATION NUMBER: PCT/US01/50423
PRIOR FILING DATE: 2001-10-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: Synthetic construct
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ORGANISM: Artificial Sequence
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CURRENT FILING DATE: 2002-04-24
PRIOR APPLICATION NUMBER: US 09/694,992
PRIOR FILING DATE: 2000-10-24
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NUMBER OF SEQ ID NOS: 11
SOFTWARE: PatentIn version 3.1
SEQ ID NO 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: STABILIZATION OF RADIOPHARMACEUTICAL COMPOSITIONS FILE REFERENCE: 09744-016001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Cyr, John E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 equence 5, Application US/10131543
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local
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9; Conserv
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100.0%; Pred. No. 0.
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0.0026;
es 0; Indels
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CURRENT FILING DATE: 2002-04-24
PRIOR APPLICATION NUMBER: US 09/695,494
PRIOR FILING DATE: 2000-10-24
PRIOR PILING DATE: 2001-10-24
PRIOR FILING DATE: 2011-10-24
NUMBER OF SEQ ID NOS: 29
SEQ ID NOS: 29
SEQ ID NOS 5
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LOCATION: 30
US-10-131-546-5
APPLICANT: KASET, MATTHEW R.
TITLE OF INVENTION: GENES EXPRESSED IN TREATED HUMAN C3A LIVER CELL CULTURES FILE REFERENCE: PA-0035 US
CURRENT APPLICATION NUMBER: US/09/919,039
CURRENT FILING DATE: 2002-09-09
PRIOR APPLICATION NUMBER: 60/222,113
NUMBER OF SEQ ID NOS: 401
                                                                                                                                                                                                                                             Sequence 33, Applica Publication No. US20 GENERAL INFORMATION:
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Best Local S
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; LOCATION: 30
US-10-131-346-5
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Publication No.
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Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Cyr, John E.
APPLICANT: Pearson, Daniel A.
TITLE OF INVENTION: STABILIZATION OF RADIOPHARMACEUTICAL COMPOSITIONS
TITLE OF INVENTION: USING HYDROPHILIC THIOETHERS AND HYDROPHILIC 6-HYDROXY
TITLE OF INVENTION: CHROMANS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION
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LENGTH: 30
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TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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OTHER INFORMATION: Synthetic construct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence
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9; Conserv
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                                                                                                                                                                                                                                                                Application US/09919039
lo. US20030108871A1
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Pred. No. 0.0026;
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Pred. No.
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DB 9; Length 30;

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; ORGANISM: Homo Sapiens
US-10-017-724-6
                                                                                  US-10-142-935-9
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                                                                                                                                                                                                                                                                                            Sequence 9, Application US/10142935
Publication No. US2003004418A1
GENERAL INFORMATION:
APPLICANT: DAVIS, Stacey
APPLICANT: HOOK, Magnus A.O.
TITLE OF INVENTION: METHOD AND COMPOSITIONS FOR INHIBITING THROMBIN-INDUCED COAGULAT:
FILE REFERENCE: P07201US01/BAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:

APPLICANT: McCarthy, Jeanette
TITLE OF INVENTION: DIAGNOSIS AND TREATMENT OF VASCULAR DISEASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 6
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                                          Query Match
                                                                                                                                                        SEQ ID NO 9
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Best Local
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      Matches
                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/10/142,935
CURRENT FILING DATE: 2002-05-13
PRIOR APPLICATION NUMBER: US 60/290,072
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/10/017,724
CURRENT FILING DATE: 2001-12-14
PRIOR APPLICATION NUMBER: US 60/317,178
PRIOR FILING DATE: 2001-09-05
PRIOR APPLICATION NUMBER: US 60/329,958
PRIOR APPLICATION NUMBER: US 60/329,958
PRIOR FILING DATE: 2001-10-16
                                                                                                                                                                          PRIOR FILING DATE: 2001-05-13
NUMBER OF SEQ ID NOS: 11
SOFTWARE: PatentIn version 3.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
NAME/KEY: misc feature
OTHER INFORMATION: Incyte ID No. US20030108871A1 3393861CD1
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TYPE: PRT
ORGANISM: Homo sapiens
                                                                                        TYPE: PRT ORGANISM: Staphylococcus epidermidis
                                                                                                                                      LENGTH: 10
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  Local Similarity 100 hes 6; Conservative
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Similarity 100.0%;
9; Conservative (
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  70.6%; Score 36; DB 9;
100.0%; Pred. No. 0.69;
Live 0; Mismatches
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Pred. No. 0.057;
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Pred. No. 0.057;
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                                    Length 10;
  Indels
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  Gaps
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GENERAL INFORMATION: BAPFLICANT: DAVIS, Stacey

APPLICANT: HOOK, Magnus A.O.

TITLE OF INVENTION: METHOD AND COMPOSITIONS FOR

FILE REFERENCE: P07201US01/BAS

CURRENT FILING DATE: 2002-05-13

PRIOR APPLICATION NUMBER: US 60/290,072

PRIOR FILING DATE: 2001-05-13

NUMBER OF SEQ ID NOS: 11

SOFTWARE: PatentIn version 3.1

SEQ ID NO 5

LENGTH: 20

TYPE: PRT
ORGANISM: Staphylococcus epidermidis

US-10-142-935-5

TO.6%: Score 36: DE
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US-10-213-512-296
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GENERAL INFORMATION: APPLICANT: Fischer,
                               Sequence 296, A Publication No.
                                                                                                                                                                                              Matches
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Publication No. US20030044418A1
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Best Local Similarity
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APPLICANT: HOOK, Magnus A.O.
TITLE OF INVENTION: METHOD AND COMPOSITIONS FOR INHIBITING THROMBIN-INDUCED COAGULAT
FILE REFERENCE: PO7201US01/BAS
CURRENT APPLICATION NUMBER: US/10/142,935
CURRENT FILING DATE: 2002-05-13
PRIOR APPLICATION NUMBER: US 60/290,072
PRIOR FILING DATE: 2001-05-13
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TYPE: PRT
                                                                                                                                                                                                             Local Similarity
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                                              Application US/10213512
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                                 US20030110536A1
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100.0%; Pred. No.
tive 0; Mismatc
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SEQ ID NO 296
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PRIOR FILING DATE: 1998-10-22
PRIOR APPLICATION NUMBER: US 09/071,838
PRIOR FILING DATE: 1998-05-01
NUMBER OF SEC. 1998-05-01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION APPLICANT: Fisch
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CURRENT APPLICATION NUMBER: US/10/213,512
CURRENT FILING DATE: 2002-08-06
                                    TELEFAX: (415) 576-03
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Arabidopsis sp
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ITLE OF INVENTION: Fruit Development i
ILE REFERENCE: 023070-086110US
                                                                                                                                                            SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/071,838 FILING DATE: 01-MAY-1998 CLASSIFICATION: 800 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ENGTH:
                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                        REFERENCE/DOCKET NUMBER: 02
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Nucleic Acids That Control : TITLE OF INVENTION: Fruit Development in Plants
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                                                                                                                       NAME: Bastian, Kevin L.
REGISTRATION NUMBER: 34,774
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                                                                         TELEPHONE:
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5. US20020152501A1
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                                                                                                                                                                                                                                                                                                                                                                                           California
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Yadegari, Ramin
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66 amino acids
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Yadegari, Ramin
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ENTION: Nucleic Acids That Control Seed and
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                                                                     (415) 576-0200
                                                    576-0300
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85.7%;
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Center, Eighth Floor
                                                                                                           023070-086100US
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Pred. No.
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-799-875-5
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Thes 6; Conserve
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CURRENT FILING DATE: 2001-05-23
PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR PRIOR PRIOR NUMBER: US 09/632,366
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR FILING DATE: 2000-08-03
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: US 09/236,359
PRIOR PILING DATE: 2000-09-27
PRIOR PILING DATE: 2000-09-27
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SEQ ID NO 5
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Patent No. US20020034780A1
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APPLICANT:
                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT FILING DATE: 2001-05-23
                                                                                                                                                                                                                                 TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY FILE REFERENCE: Aeomica-X-1
                                                                                                                                                                                                                                                                                                                                 APPLICANT: Penn, Sharron G. APPLICANT: Rank, David R.
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PRIOR FILING DATE: 2000-09-12
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PRIOR APPLICATION NUMBER: 60/18
PRIOR FILING DATE: 2000-02-11
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CURRENT FILING DATE: 2001-03-06
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Chen, Wensheng
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Williamson, Mark
                                                                                                                                                                                                                                                                                                                               Rank, David R.
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85.7%;
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Pred. No. 1
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Pred. No.
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FILING DATE: 2001-01-30

NUMBER: PCT/US01/00667

2001-01-30

FILING DATE: 2001-01-30

APPLICATION NUMBER: PCT/US01/00669 APPLICATION NUMBER: PCT/US01/00664

2001-01-30

ILING DATE:

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US-09-350-874-32
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SEQ ID NO 34308
LENCTH: 543
TYPE: PRT
                                                                                                                                                                                                                                                                Sequence 32, Application US/09350874 Patent No. US20020019020A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
             CURRENT APPLICATION NUMBER: US/09/350,874
CURRENT FILING DATE: 1999-07-09
EARLIER APPLICATION NUMBER: USSN 60/110,277
EARLIER FILING DATE: 1998-11-30
EARLIER APPLICATION NUMBER: USSN 60/110,033
EARLIER FILING DATE: 1998-11-25
EARLIER APPLICATION NUMBER: USSN 60/109,333
EARLIER APPLICATION NUMBER: USSN 60/109,333
                                                                                                                                                                               APPLICANT: Rhodes, Kenneth
APPLICANT: An, Wengian
APPLICANT: An, Wengian
TITLE OF INVENTION: METHODS FOR TREATING CARDIOVASCULAR DISORDERS
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PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 09/608,408
PRIOR FILING DATE: 2000-06-30
PRIOR APPLICATION NUMBER: US 09/774,203
PRIOR FILING DATE: 2001-01-29
NUMBER OF SEQ ID NOS: 49117
                                                                                                                                                                       FILE REFERENCE: MNI-069
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
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APPLICATION NUMBER: PCT/US01/00670
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APPLICATION NUMBER: PCT/US01/00662
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NUMBER: USSN 09/298,731
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IN ADULT LIVER, SIGNAL = 1.9
IN HEART, SIGNAL = 1.4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 34; DB 10;
Pred. No. 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    N HBL100,
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BRAIN, SIGNAL = 3.3
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HELA, SIGNAL = 1.
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AI002818.1, EVALUE 5.00e-24
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; EARLIER FILING DATE: 1999-04-23; NUMBER OF SEQ ID NOS: 72; SOFTWARE: PATENTIN Ver. 2.0; SEQ ID NO 32; LENGTH: 256; TYPE: PRT; ORGANISM: Homo sapiens US-09-350-874-32
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                                        Query Match
Best Local S
Matches 6
20 GHTPLSKK 27
                                        l Similarity
6; Conserv
                   1 GHRPLDKK
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75.0%;
                                         Score 34; DB Pred. No. 61; 0; Mismatches
                                                               DB
                                                               10;
                                         2;
                                                              Length 256;
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Search completed: June 27, Job time: 28.9 secs

2003, 17:30:22

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Minimum
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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                 Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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DB seq length:
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                 of hits satisfying chosen parameters:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Match
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51
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*
/cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
/cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
/cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
/cgn2_6/ptodata/1/iaa/BCTUS_COMB.pep:*
/cgn2_6/ptodata/1/iaa/backfiles1.pep:*
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                                                                    US-08-468-964B-10
US-07-871-82A-10
US-08-253-678A-10
US-08-253-678A-10
US-08-253-678A-10
US-08-253-578A-10
US-08-290-853-35
US-08-266-178A-9
US-08-268-699-11
US-08-058-699-11
US-09-399-313-32
US-09-399-313-32
US-09-399-313-32
US-09-315-944-25-3
US-08-107-684B-9
US-08-107-684B-9
US-08-227-372-1
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                                               US-09-134-001C-4644
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Sequence 1, Appli Sequence 12, Appl Sequence 10, Appl Sequence 35, Appl Sequence 35, Appl Sequence 9, Appli Sequence 9, Appli Sequence 9, Appli Sequence 10, Appl Sequence 10, Appl Sequence 11, Appl Sequence 11, Appl Sequence 11, Appl Sequence 12, Appl Sequence 32, Appl Sequence 9, Appli Sequence 9, Appli Sequence 12, Appl Sequence 13, Appl Sequence 13, Appl Sequence 13, Appl Sequence 14, Appl Sequence 14, Appl Sequence 16, Appl Sequence 17, Appl Sequence 17, Appl Sequence 18, Appl Sequence 16, Appl Sequence 17, Appl Sequence 17, Appl Sequence 18, Appl Sequence 18,
                                                                          Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Description
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US-08-448-547-1
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Patent No.
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US-09-241-333-1	US-08-990-114-1	US-09-163-444-2	US-09-137-440-50	US-08-833-485-50	US-08-306-063-50	US-08-476-008-50	US-08-979-917A-4	US-08-558-823-19	US-09-227-357-348	US-08-976-838-15	US-08-525-940-15	US-08-368-852-15	US-09-180-109A-31	US-09-180-109A-35	US-09-199-637A-33	US-09-227-357-357	US-08-507-124-4
Sequence 1, Appli	Sequence 1, Appli	Sequence 2, Appli	Sequence 50, Appl	Sequence 50, Appl	•	Sequence 50, Appl	Sequence 4, Appli	Sequence 19, Appl	œ	Sequence 15, Appl	•	Sequence 15, Appl	Sequence 31, Appl	•	Sequence 33, Appl	Sequence 357, App	Sequence 4, Appli

ALIGNMENTS

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; MOLECULE TYPE: peptide US-08-448-547-1
                                                                                                                                                                                                                                                             FILING DATE: 30-JUN-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/JP
FILING DATE: 01-NOV-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 5-2
FILING DATE: 02-NOV-1993
ATTORNEY_AGENT INFORMATION:
NAME: Nakamura, Dean H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                   SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
TYPE: amino acid
STRANDEDNESS:
TOPOLOGY: linear
                                                                                                                                           INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 20037
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                          NAME: Nakamura, Dean H.
REGISTRATION NUMBER: 33,981
REFERENCE/DOCKET NUMBER: Q-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)293-7060
TELEPHONE: (202)293-7060
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: ZIVU
STREET: Washington
STATE: D.C.
STATE: D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Ito, Yumiko
TITLE OF INVENTION: ANTIGHUMAN SOLUBLE FEBRIN ANTIBODY,
TITLE OF INVENTION: HYBRIDOMA, AND IMMUNOASSAYING METHOD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                                TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/448,547
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1, Application US/08448547
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                E: SUGHRUE, MION, ZINN, MACPEAK & SEAS 2100 Pennsylvania Avenue, N.W.
                                                                                                   10 amino acids
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Kohno, Isao
Inuzuka, Kimiko
                                                                                                                                                            (202) 293-7860
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밁 8

Matches

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US-08-470-152-12
                                                                                                                                                                                                                                                                                                                             ; MOLECULE TYPE: peptide US-08-486-135-12
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                                                                                                    Sequence 12, Application US/08470152 Patent No. 5780007
                                                                                    Patent No. 5780007
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 12
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                                                                                                                                                                                                                                                                                                                                                                                                                        TELEX: 910-221-5317
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 28 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: NO. 5720934nan, Kevin
REGISTRATION NUMBER: 35,303
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: LEM PC compatible
ODERATING SYSTEM: PC-DOS/MS-DOS
                                                       APPLICANT:
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TITLE OF INVENTION: Imaging
NUMBER OF SEQUENCES: 19
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APPLICANT: Civitello,
                                                                                                                                                                                                                                                                              Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: FILING DATE: 06-JUN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE:
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9; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       E: Banner & Allegretti, Ltd.
10 South Wacker Drive, Suite 3000
   Lister-James, John
Civitello, Edgar R
                                           Buttram, Scott
                                   McBride, William
                                                                                                                                                                                                                                                                                                                                                                                                                                                        312-715-1234
                                                                                                                                                                                                                                                     100.0%; Score 51; DB 1; Length 28; llarity 100.0%; Pred. No. 0.0018; Conservative 0; Mismatches 0; Indels
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                                                                   Dean, Richard T
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McBride, William
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Pred. No.
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Best Local (
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INFORMATION FOR SEQ ID NO:
COMPUTER READABLE FORM.

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: BM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
CURRENT APPLICATION NUMBER: US/08/468,964B
                                                                                                   ZIP: 03053
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                             TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
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LENGTH: 28 amino acid:
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NAME: NO. 5780007/nan, Kevin
REGISTRATION NUMBER: 35,303
REFERENCE/DOCKET NUMBER: 92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOPTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/470,152
FILING DATE: 06-UN-1995
CLASSIFICATION: 424
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TELEPHONE: 312-715-1000
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TITLE OF INVENTION: Im.
NUMBER OF SEQUENCES: 1:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                        STATE:
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                                                                                                                                                     COUNTRY:
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OPERATING SYSTEM:
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CITY: Chicago
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9; Conserva
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amino acid
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                                                                                                                                                                                                                                                                                                                             Buttram, Scott
McBride, William
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Civitello, Edgar R
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10 South Wacker Drive, Suite 3000
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IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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Length 28; Indels

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APPLICANT: Dean, Richard T.

APPLICANT: Buttram, Scott

APPLICANT: McBride, William
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                                                                                                                                                                                         NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Release #1.0, VACURENT APPLICATION DATA:
APPLICATION UMBER: US/07/871,282A
FILING DATE: 20-APR-1992
CLASSIFICATION: 424
                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 28 amino acids
TYPE: amino acid
STRANDEDNESS: not relev
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LENGTH: 28 amino acids
                                                                                            TOPOLOGY: 1: MOLECULE TYPE:
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TITLE OF INVENTION:
NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
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Local Similarity 100.0%;
hes 9; Conservative 0
                                                                                                                                                                                                                                                      NAME: McDaniels, Patricia A.
REGISTRATION NUMBER: 33,194
REFERENCE/DOCKET NUMBER: DITI 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: y L.
                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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STRANDEDNESS: not
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Civitello, Edgar R.
IVENTION: Technetium-99m Labeled Peptides for
              100.0%; ilarity 100.0%; Conservative 0
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             Score 51; DB 2; I Pred. No. 0.0018; 0; Mismatches 0;
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Pred. No. 0.0018;
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                                            DB 2; Length 28;
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RESULT 7
US-08-253-678A-10
: Sequence 10, Application US/08253678A
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TELEX: 910-221-5317
INFORMATION FOR SEQ ID NO: 3
SEQUENCE CHARACTERISTICS:
LENGTH: 28 amino acids
                                                                                                                                                     Matches
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                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                              OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TORNEY/AGENT TORNEY/AGENT NAME: NO. 5989519nan, Kevin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/08/290,853
FILING DATE: 11-OCT-1994
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION:
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STATE: Illinois
                                                                                                                                                                                                                                                                                         NAME/KEY: Modified-site
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lo. 598951!
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10 South Wacker Drive, Suite 3000
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Imaging Inflammation
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radiolabel binding moiety."
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/note= "The carboxyl terminus is modified
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                                                                                                                                                                Score 51; DB 2;
Pred. No. 0.0018;
                                                                                                                                                     Mismatches
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                           Patent No.
                                                                                                                                                                                                                                                      GENERAL INFORMATION:
APPLICANT: Dean, I
APPLICANT: Buttra
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INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
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OMPUTER READABLE FORM:
"MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                             NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
ADDRESSEE: Diatide, Inc.
STREET: 9 Delta Drive
                                                                                                                                                                           TITLE OF INVENTION: TECHNET INTLE OF INVENTION: IMAGING
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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ADDRESSEE: Diatide, Inc.
STREET: 9 Delta Drive
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/08/253,678A FILING DATE: 03-JUN-1994 CLASSIFICATION: 424
                                                                  COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: DITI 112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: McDaniels, Patricia A. REGISTRATION NUMBER: 33,194
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH:
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McBride, William
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/ENTION: TECHNETIUM-99m LABELED PEPTIDES FOR
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McBride, William
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US-08-170-299-10
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                    TELEFAX: 312-715-1234
TELEX: 910-221-5317
INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                           COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC COMPUTER: PC-DOS/MS-DOS
SOFTMARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/170,299
FILLING DATE: 09-EBB-1995
CLASSIFICATION: 514
CCLASSIFICATION: 514
                             SEQUENCE CHARACTERISTICS:
LENGTH: 28 amino acids
                                                                                                             TELECOMMUNICATION INFORMATION: TELEPHONE: 312-715-1000
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TOPOLOGY: linear
MOLECULE TYPE: peptide
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (603) 437-8970
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CURRENT APPLICATION DATA:
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                                                                              TELEPHONE: 312-715-1234
                                                                                                                                              NAME: No. 6086849nan, Kevin
REGISTRATION NUMBER: 35,303
REFERENCE/DOCKET NUMBER: 92
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                               28 amino acids
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10 South Wacker Drive, Suite 3000
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McBride, William
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Technetium-99m Labeled Peptides for
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n Release #1.0, Version #1.30
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Best Local
                                                                 Query Match
Best Local
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COMPUTER: IBM PC COMPATIBLE

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/290,853

FILING DATE: 11-OCT-1994

CLASSIFICATION: 424

ATTORNEY/AGENT INFORMATION:

NAME: NO. 5989519nan, Kevin E

REGISTRATION NUMBER: 35,303

REFERENCE/DOCKET NUMBER: 92,112-H
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SEQUENCE CHARACTERISTICS:
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TELEPHONE: 312-715-1000
TELEFAX: 312-715-1234
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MEDIUM TYPE: Floppy
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APPLICANT: Buttram, Scott
                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: peptide
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                                                                                                                            OTHER INFORMATION:
OTHER INFORMATION:
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OTHER INFORMATION:
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TELEFAX: 910-221-5317
TELEFAX: 910-221-5317
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                                                                                                                                                                                           NAME/KEY: Modified-site
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                                                                                                                                                                           LOCATION:
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                                                                Similarity
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: Illinois
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10 South Wacker Drive, Suite 3000
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Imaging Inflammation 
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/note= "
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/note= menor terminal residue is
/rote-2-carbonyl, the thiol of the
pyridime-2-carbonyl, the thiol group.
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                                                              Score 51; DB 2;
Pred. No. 0.0019;
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Pred. No. 0.0018;
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                                               Mismatches
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                                                                                                          Sequence 9, Application US/08484774 Patent No. 5807538
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Patent No. 571193
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Best Local
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APPLICANT: Dean, Richard T
APPLICANT: Lees, Robert S.
                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: 312-715-1234
TELEX: 910-221-5317
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                           APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/472,535
FILING DATE: 07-UN-1995
CLASSIFICATION: 424
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: LUCITY: Chicago
CITY: Illinois
TTATE: ITSA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: NO. 5711931nan, Kevin
REGISTRATION NUMBER: 35,303
REFERENCE/DOCKET NUMBER: 92.
               TITLE OF INVENTION:
 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION: TELEPHONE: 312-715-1000
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Buttram, Scott
TITLE OF INVENTION: Technetium-99m Label
TITLE OF INVENTION: Imaging Inflammation
                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: Modified LOCATION: 1..3 OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE:
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                                                                                                                                                                                                                                          1 GHRPLDKKR 9
                                                                                                                                                                                                                                                                                       Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       amino acid
XGY: linear
                                                                                                                                                                                                          GHRPLDKKR 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       31 amino acids
                                             Buttram,
                                                          Dean, Richard T
Lees, Robert S.
                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               E: Floppy disk
IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                        Modified-site
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lmaging
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             Technetium-99m Labeled Peptides Imaging Inflammation
                                               Scott
                                                                                                                                                                                                                                                                                                                                                      /note= "The amino terminal residue is pyridine-2-carbonyl; the thiol of the cysteine residue is protected by an acetoamidomethyl
                                                                                                                                                                                                                                                                                                                                                                                                        /label= Picolinoyl
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Pred. No. 0.002;
                                                                                                                                                                                                                                                                           Mismatches
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Length 31 Indels

0

Gaps

COUNTRY:

60606

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GENERAL INFORMATION:
                                        MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
GURRENT APPLICATION DATA:
                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/08/484
FILING DATE: 07-UUN-1995
CLASSIFICATION: 424
ATTORNEY AGENT INFORMATION:
NAME: NO. 5807538nan, Kevin E
REGISTRATION NUMBER: 35,03
REFERENCE/DOCKET NUMBER: 92,1
                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: 312-715-100
TELEPAX: 312-715-1234
TELEX: 910-221-5317
INFORMATION FOR SEQ ID NO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 31 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                           CITY: Chicago
STATE: Illinois
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION:
OTHER INFORMATION:
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-715-1000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Release #1.0, Version #1.25
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ADDRESSEE: Banner & Allegretti, Ltd.
STREET: 10 South Wacker Drive, Suite 3000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: Modified-site
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5. 6017510
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9; Conserv
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10 South Wacker Drive, Suite 3000
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: Illinois
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                                   US/08/266,178A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /label= Picolinoyl
/note= "The amino terminal residue is
pyridine-2-carbonyl; the thiol of the cysteine
residue is protected by an acetoamidomethyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US/08/484,774
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 51;
Pred. No.
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                                                                                  #1.25
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                                                              TELEFAX: 206-548-2329
INFORMATION FOR SEQ ID NO:
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                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Garner, Ian
APPLICANT: Dalrymple, Micha
APPLICANT: Prunkard, Donna
APPLICANT: Foster, Donald C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local
                                                                                                                                REFERENCE/DOCKET NUMBER: 9:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patenth Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
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TELEX: 910-221-5317
INFORMATION FOR SEQ ID NO.
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
                                                                                                                 TELEPHONE:
                                                                                                                                                                                  NAME: Parker, Gary E
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                   FILING DATE:
CLASSIFICATION:
                         ENGTH:
                                                                                                                                                                                                                                                                                              APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET:
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FEATURE:
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OTHER INFORMATION: /label= Picolinoyl
OTHER INFORMATION: /note= "The amino of the information of the information
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TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
TOPOLOGY: linear
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REGISTRATION NUMBER: 35,303
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    98105
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amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WA
               491 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          E: ZymoGenetics, Inc.
4225 Roosevelt Way, N.E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                USA
                                                                                                    206-547-8080 ext 322
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Foster, Donald C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     31 amino acids
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                                                                                                                                                                   31-648
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                                                              4:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Michael A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note= "The amino terminal residue is
pyridine-2-carbonyl; the thiol of the cysteine
residue is protected by an acetoamidomethyl
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Query Match Best Local Matches

US-08-266-178A-9

Sequence 9, Patent No.

APPLICANT:

COUNTRY:

USA

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STREET:

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Search completed: June 27, 2003, 17:19:04 Job time : 10.9 secs
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                                                                                                                                                                                                                                                                                                                                                                                                                   ; TYPE: AMINO ACID
; STRANDEDNESS:
; TOPOLOGY: Linear
US-08-058-699-10
                                                                                                                                                                                                                                                                                                          Query Match
Best Local S
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Best Local S
                                                                                                                                                                                                                                                                                Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PS/2 Model 50Z or 55SX
OPERATING SYSTEM: MS-DOS (Version 5.0)
SOFTWARE: WordPerfect (Version 5.1)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/058,699
FILING DATE: 1930503
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: N/A
APPLICATION NUMBER: N/A
APPLICATION NUMBER: N/A
APPLICATION DATA:
APPLICATION NUMBER: N/A
APPLICATION NUM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Janis K. Fraser, Ph.D.
REGISTRATION NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 054
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: U.S.A.
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Edgar Haber
APPLICANT: Christoph Bode
APPLICANT: Marschall S. Runge
TITLE OF INVENTION: FIBRIN-TARGETED INHIBITORS OF
TITLE OF INVENTION: THROMBIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: \u_
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45 GHRPLDKKR 53
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                                                                                                                                                                                                                                                                     h 80.4%; Score 41; DB 1; Similarity 100.0%; Pred. No. 1.9e+(7; Conservative 0; Mismatches
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Result
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 Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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1 GPRVVERHQSA 11
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2: pir2:*
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ОУ	Query Matc Best Local Matches	RESULT 2 XASNPC angiotensin-converting enzyme: C;Species: Vipera aspis (aspic C;Date: 31-Mar-1993 #sequence_) C;Accession: A60377 R;Komori, Y: Sugihara, H. Int. J. Biochem. 22, 767-771, 1 A;Title: Characterization of a A;Reference number: A60377 A;Molecule type: protein A;Residues: 1-10 <kom> C;Superfamily: bradykinin-poter C;Keywords: angiotensin-convert F;1/Modified site: pyrrolidone</kom>	Best Local Matches Qy	RESULT 1 JN0440 peptide-N4 (N-acetyl-beta-glucosaminyl)and N; Alternate names: glycopeptidase C; Species: Prunus dulcis var. sativa (sw. C; Date: 30-Sep-193) #sequence_revision 30 C; Accession: JN0440 R; Kaliberda, E.I.; Shemyakin, V.V.; Anton Bioorg. Khim. 16, 751-757, 1990 A; Title: Purification of sweet almond gl) A; Reference number: JN0440 A; Accession: JN0440 A; Accession: JN0440 A; Molecule type: protein A; Regidues: 1-10 < KAL> C; Keywords: hydrolase	
1 GPRV : 5 GPKV	h Simi 3;	Vipera Vipera Vipera A-Mar-1 A	Sim 4; 2 PR: 1 1	N4-(N-ace nate names se Prunus 30-Sep-19 30-Sep-19 30-Sep-19 Midn. 10-19 Khim. 16.6 Khim. 16.6 Khim. 18.7 Khim. 16.7 Khim. 16.7 Khim. 17.0 Purifice nee numbe pion: JN04 Jle type: les: hydrods: hydrods:	
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RESULT 3

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A;Residues.
C;Keywords:
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C;Species: Carcharhinus plumbeus (sandbar shark)
C;Date: 11-Aug-1995 #sequence_revision 11-Aug-1995 #text_change 20-Jun-2000
C;Accession: B56899
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                                                                                                                                                                Comp. Biochem. Physiol. B 103, 563-568, 1992
A;Title: Purification of a novel heterodimer from shark
A;Reference number: A56899; MUID:93092592; PMID:1458832
A;Accession: B56899
                                                                                                                                                                                                                           R; Vazquez-Moreno, L.; Porath, J.;
                                                                                                                                                                                                                                                                                                           B56899
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A;Title: Organization of the gene (RHCE) encoding the human blood group RhCcBe antigens A;Reference number: I54193; MUID:94245182; PMID:8188244
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A; Residues: 1-11 < RES>
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C;Accession: I54193
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C;Species: Homo sapiens (man)
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R;Kawakami, T.; Kamc
submitted to JIPID,
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C;Species: Oryza sativa (rice)
                                                                                                                  ;Molecule type: protein ;Residues: 1-10 <VAZ>
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ber: PC4267
                                       Conservative
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                                                  31.6%;
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Pred. No. 2e+0
0; Mismatches
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Pred. No. 2
                                                                                                                                                                                                          Schluter, S.F.; Marchalonis, J.J
3-568, 1992
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A;Molecule type: protein
A;Residues: 1-11 <FOR>
C;Keywords: glycolysis; phosphotransferase
                                                                                                                              A;Reference number: A14454; MUID:80004524; PMID:157899
A;Accession: A14454
                                                                                                                                                                          Biochem. Soc. Trans. 7, 721-72
A; Title: The N-terminal amino
                                                                                                                                                                                                                                            C;Species: Ovis orientalis aries, Ovis ammon aries (domestic sheep)
C;Date: 05-Jun-1987 #sequence_revision 05-Jun-1987 #text_change 28-Apr-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Molecule type: DNA
A;Residues: 1-11 <YAM>
A;Esidues: 1-11 <YAM>
A;Experimental source: B lymphocyte
C;Keywords: heterotetramer; immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RyYamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G. J. Exp. Med. 173, 395-407, 1991
A;Title: Preferential utilization of specific immunoglobulin heavy chain diversity A;Reference number: PT0222; MUID:91108337; PMID:1899102
A;Accession: PT0302
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ig heavy chain CRD3 region (clone 5-112) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 30-Sep_1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996
                                                                                                                                                                                                                  R; Fordyce,
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                                                                                                                                                                                                                                                                                          6-phosphofructokinase (EC 2.7.1.11) -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 33K protein 3218 - rice (strain Nohonbare) (rragment)
C;Species: Oryza sativa (rice)
C;Date: 03-Feb-1994 #sequence_revision 03-Feb-1994 #text_change 11-Apr-1995
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A;Residues: 1-11 <TSU5-
A;Experimental source: bran
C;Comment: molecular weight 33K,
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Pred. No. 3.2e
2; Mismatches
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Pred. No: 5e+03;
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ERHOSA 11

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C;Accession: A37765
R;Theroux, S.J.; Redlinger, T.E.; Fuller, R.J. Bacteriol. 172, 4497-4504, 1990
A;Title: Gene encoding the 5.7-kilodalton chi A;Reference number: A37765; MUID:90330558; PW A;Accession: A37765
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Molecule type: DNA
A;Residues: 1-6 <THE>
A;Cross-references: GB:M33964
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C;Species: Chloroflexus aurantiacus
C;Date: 31-May-1991 #text_change
C;Accession: A37765
C;Accession: A37765
R;Theroux, S.J.; Redlinger, T.E.; Fuller, R.C.; Robinson, S.J.
                 Ś
                                                                                                                                                                      R;Carraway, R.E.; Cochrane, D.E.; Boucher, W.; Mitra, S.P.
J. Immunol. 143, 1680-1684, 1989
A;Tille: Structures of histamine-releasing peptides formed
A;Reference number: A45800; MUID:89341406; PMID:2474609
                                                                                                                                                                                                                                              serum albumin - dog (fragment)
C;Species: Canis lupus familiaris (dog)
C;Date: 03-Jun-1993 #sequence_revision
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A;Title: Amino acid sequence of CAP(2b), an insect cardioacceleratory peptide A;Reference number: S66646; MUID:96013159; PMID:7556618
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cardioacceleratory protein 2b - tobacco hornworm C; Species: Manduca sexta (tobacco hornworm)
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Pred. No. 2.8e+05;
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probable antigen 5 - Mycobacterium leprae (fragment)
(;Species: Mycobacterium leprae
(;Date: 10-Apr-1992 #sequence_revision 10-Apr-1992 #text_change 18-Jun-1993
(;Accession: E41476
R;Hartskeerl, R.A.; van Rens, R.M.; Stabel, L.F.E.M.; de Wit, M.Y.L.; Klatser, Infect. Immun. 58, 2821-2827, 1990
A;Title: Selection and characterization of recombinant clones that produce Myco A;Reference number: A41476; MUID:90354041; PMID:1696931
A;Accession: E41476
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          R;Roessner, C.A.; Warren, M.J.; Santander, P.J.; Atshaves, B.P.; Ozaki, S.; Stolowich, FEBS Lett. 301, 73-78, 1992
A;Title: Expression of 9 Salmonella typhimurium enzymes for cobinamide synthesis. Iden: A;Reference number: S21127; MUID:93083628; PMID:1451790
A;Accession: S21127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              precorrin methyltransferase - Salmonella typhimurium
C;Species: Salmonella typhimurium
C;Date: 22-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 10-Nov-1995
C;Accession: S21127
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Exp. Hematol. 19, 704-709, 1991
A;Title: Purification and amino acid analysis of a human macrophage cytotoxicity-induci A;Reference number: A60427; MUID:91372335; PMID:1909970
A;Accession: A60427
                                                                                         A;Status: preliminary; not compared with conceptual translation A;Molecule type: DNA A;Residues: 1-11 <HAR>
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A;Molecule type: protein
A;Residues: 1-11 <ROE>
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C;Species: Homo sapiens (man)
C;Date: 12-Feb-1993 #sequence_revision 12-Feb-1993 #text_change 18-Jun-1993
C;Accession: A66427
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5 PRV 7

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Page 4
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RESULT 15
PS0259
39K protein 3225 - rice (strain Nihonbare) (fragment)
C;Species: Oryza sativa (rice)
R;Tsugita, A;Farcles (rice)
Search completed: June 27, 2003, 17:34:17
Search completed: June 27, 2003, 17:34:17
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Result
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Maximum Match 100%
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   protein search, using sw model
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57
1 GPRVVERHQSA 11
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Gapop 10.0 , Gapext 0.
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UF03 MOUSES
UPA5 HUMAN
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ALL7 LOCMI
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P04277 homo sapien
P32080 homo sapien
P38641 mus musculu
P30091 homo sapien
P17339 morganella
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P83564
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P31351 vipera aspi
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4 leptinotars
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P02729	P38642	. P36960	P31890	P12997	P01858	P81350	P28499	P82684	P30424	P30423	P09037
homo sapien	mus musculu	carnobacter	ascaris suu	citrobacter	homo sapien	clostridium	oncorhynchu	carausius m	bothrops in	bothrops in	crinia geor

ALIGNMENTS

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RESULT 3
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P43172;
01-NOV-1995 (Rel. 3
01-NOV-1995 (Rel. 3
01-FEB-1996 (Rel. 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Komori Y., Sugihara H.;
"Characterization of a new inhibitor for angiotensin "Characterization of a new inhibitor for angiotensin enzyme from the venom of Vipera aspis aspis.";
Int. J. Biochem. 22:767-771(1990)
Int. J. Biochem. 22:767-771(1990)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Lepidosauria; Squamata; Scleroglossa; Serpentes; Colubroidea; Viperidae; Viperinae; Vipera.
NCBI_TaxID=8706;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   enzyme inhibitor).
Vipera aspis (Aspic viper)
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01-FEB-1994 (Rel.
01-FEB-1994 (Rel.
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Neuropeptide;
                                                                                                                         MEDLINE=95380362; PubMed=7651904;
Cowden C., Stretton A.O.W.;
"Eight novel FMRFamide-like neuro
                                                                                                                                                                                                                         NCBI_TaxID=6253;
                                                                                                                                                                                                                                                                                          FMRFamide-like neuropeptide AF9 Ascaris suum (Pig roundworm) (A
                                                                                                                                                                                                                                                                                                                                                                                                                                                ASCSU
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                                                                                  Peptides 16:491-500(1995)
                                                                                                   Ascaris suum.";
                                                                                                                                                                                                                                                       Ascarididae;
                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Nematoda; Chromadorea; Ascaridida; Ascaridoidea;
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                                                                 SIMILARITY:
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33, Last annotation update)
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3BA827C327686773
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                                                                                               5_HUMAN
URA6_HUMAN
P32080;
01-OCT-1993
                                                                                                                                                                                                                                                                                                                                                                                                                    Mogard M.H., Kobayashi R., Chen C.F., Le Shively J.E., Walsh J.H.;
"The amino acid sequence of kinetensin, pepsin-treated human plasma: homology wineurotensin and angiotensin.";
Biochem. Biophys. Res. Commun. 136:983-
                                                                                                                                                                                                                                                                                                                     SPECIES=Human, Bovine, and Rabbit;
MEDLINE=87194805; PubMed=2437111;
Carraway R.E., Mitra S.P., Cochrane D.E.;
"Structure of a biologically active neurotensin-related obtained from pepsin-treated albumin(s).";
J. Biol. Chem. 262:5968-5973(1987).
-!- FUNCTION: REGULATION OF FAT DIGESTION, LIPID ABSORPT
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P04277;
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                                Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                             01-OCT-1993 (Rel. 27, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Unknown protein from 2D-page of red blood cells
                                                                                                                                                                                                                                                                                       PIR; A03239; ABHUSK.
PIR; A26693; A26693.
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MEDLINE=86242180; PubMed=3087352;
Vohavashi R., Chen C.
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Mammalia; Eutheria; Primates;
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15-JUN-2002 (Rel. 41, Last annotation update)
Neurotensin-related peptide (NRP) (Kinetensin)
SEQUENCE
                 NCBI_TaxID=9606;
                                                     Homo sapiens (Human)
                                                                                                                                                                                                                                                                   SEQUENCE
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100.0%; Pr
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0; Mismatches
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                                                                                                                                                                                                                                                                    C804DB4761F4140D CRC64;
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Pred. No. 1.1e+05
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                                Craniata; Vertebrata; Catarrhini; Hominidae;
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TISSUE=Plasma;
MEDLINE=93092937; J
Hughes G.J., Fruti
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE=95009907; PubMed=7523108; Merrick B.A., Patterson R.M., Withter L.I., He C., Sel "Separation and sequencing of familiar and novel murir using preparative two-dimensional gel electrophoresis. Electrophoresis 15:745745(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "Plasma and red blood cell protein maps: update Electrophoresis 14:1223-1231(1993).
-!- MISCELLANEOUS: ON THE 2D-GEL THE DETERMINED PROTEIN IS: 5.4, ITS MW IS: 65 KDa.
SWISS_2DPAGE, P32080; HUMAN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pasquali C., Sancne, ...
Pasquali C., Sancne, ...
Palant L., Hochstrasser D.F.;
Palant L., Hochstrasser Derivation of the coll pr
                                                                                                                                                                                                                                                                UPA5_HUMAN
P30091;
                                                                                                                                                                                                                                                                                                         HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Veri
Mammalia; Eutheria; Rodentia; Sciurognathi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-OCT 1994 (Rel. 30, Created)
01-OCT-1994 (Rel. 30, Last sequence update)
01-FEB-1995 (Rel. 31, Last annotation update)
Unknown protein from 2D-page of fibroblasts (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOUSE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Golaz O., Hughes G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=94147970;
                                                                                                                            Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                               01-APR-1993 (Rel. 25, Created
01-APR-1993 (Rel. 25, Last so
16-OCT-2001 (Rel. 40, Last ar
Unknown protein from 2D-page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TISSUE=Fibroblast;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          P38641;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TISSUE=Erythrocyte;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE
                                                                     SEQUENCE
                                                                                                     NCBI_TaxID=9606;
                                                                                                                                                                   Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=10090;
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2; Conserv
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ghes G.J., Frutiger S., Paque
Sanchez J.-C., Tissot J.-D.,
          Frutiger
                                                                                                                                                                                                                                                                                                                                                                                       4
                                                                                                                                                                                                                                                                                   STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STANDARD;
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                              PubMed=1459097
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25, Last seq
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Primates;
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Last annotation update)
2D-page of plasma (Spot
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          Paquet
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           6AA72B1DDB1B1180
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Pred. No. 1.1e
1; Mismatches
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Pred.
                                                                                                                          Craniata; Vertebrata; Catarrhini; Hominidae
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          Ravier
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                                                                                                                            Hominidae; Homo
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Appel R.D., W
      F .
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          Pasquali
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                                                                                                                                              Euteleostomi;
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; Murinae; Mus
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Best Local S
Matches 2
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Best Local
                           - AGKHP
- BPP AGKHP
P04562;
13-AUG-1987
01-FEB-1994
16-OCT-2001
  16-OCT-2001 (Rel. 28, Last sequence up
Bradykinin-potentiating peptide (Angio
enzyme inhibitor).
                                                                                                                                                                                        Hydrolase.
NON TER
SEQUENCE
                                                                                                                                                                                                                                                                                                                       Morganella.
NCBI_TaxID=582;
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amidohydrolase)
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P17339;
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                                                                                                                                                                                                                  -!- SIMILARITY: BELC

-TR: C35389; C35389.
                                                                                                                                                                                                                                                                                     MEDLINE=90264298; PubMed=2345135; Hu L.-T., Nicholson E.B., Jones B.
                                                                                                                                                                                                                                                                                                         SEQUENCE
                                                                                                                                                                                                                                                                                                                                             Morganella morganii (Proteus morganii)
Bacteria; Proteobacteria; gamma subdiv
                                                                                                                                                                                                                                                                                                                                                                UREA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE
Agkistrodon
                                                                                                                                                                                                                                                                            "Morganella morganii urease: purification,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  -!- MISCELLANEOUS: ON THE 2D-GEL THE PROTEIN IS: 4.85, ITS MW IS: 40 k
-!- MISCELLANEOUS: THIS SPOT IS ON A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sanchez J.-C., James Hochstrasser D.F.;
                                                                                                                                                                                                                                               isolation of gene sequences.";
J. Bacteriol. 172:3073-3080(1990)
-i- CATALYTIC ACTIVITY: Urea + H(
                                                                                                                                                                                                                                                                                                                                                                                            15-JUL-1999
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                                                                                                                                                   Similarity 3; Conserv
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na subunit (EC 3.5)
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                                                                 STANDARD;
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pallas
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1171
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50.0%;
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(Chinese water mocassin) (Gloydius halys
                                                                                                                                                                                                                                                                                                                                                                                 annotation update)
.5.1.5) (6 kDa subunit)
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GAMMA)(3) (BY SIM)
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Pred. No. 5.8e
2; Mismatches
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Pred. No.
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/FTId=VAR_000002
                                                                  PRT;
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                  (Angiotensin-converting
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on update)
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A POSITION THOUGHT TO
DOES NOT CORRESPOND T
                                                                                                                                                                                                                            = CO(2) + 2 NH(3).
(BY SIMILARITY).
GAMMA SUBUNIT FAMILY
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5.8e+03;
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5.8e+03;
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                                                                                                                                                                                        CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CRC64;
                                                                                                                                                                      Length 10;
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Best Local S
Matches 2
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21-JUL-1986
                                                                                                                                                                                                                                                                                                              SPECIES=A.elegantissima, and H.attenuata; Schaller H.C., Bodenmuller H.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Anthopleura elegantissima (Sea anemone), and Hydra attenuata (Hydra) (Hydra vulgaris).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID=9606, 10116, 9913, 6110, 6087;
                           REDLINE=90059923; PubMed=2583101;
Schaller H.C., Druffel-Augustin S
                                                                                                                                                                MEDLINE=82050803;
Birr C., Zachmann
                                                                                                                                                                                                               SYNTHESIS
                                                                                                                                                                                                                                                                                                "Isolation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SPECIES=Human, Rat, and Bovine; MEDLINE=82035850; PubMed=7290191;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SPECIES=Human,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rattus norvegicus (R
Bos taurus (Bovine),
                                                                                                                                                                                                                                                      Proc. Natl.
                                                                                                                                                                                                                                                                              hydra."
                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                           from coelenterates to hum
Nature 293:579-580(1981).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bodenmuller H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE
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21-JUL-1986 (Rel.
01-OCT-1996 (Rel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=86177022; PubMed=3008123;
Chi C.-W., Wang S.-Z., Xu L.-G., Wang M.-Y., Lo S.-S., Huang W.-D.;
Chi C.-W., Wang S.-Z., Xu L.-G., Wang M.-Y., Lo S.-S., Huang W.-D.;
"Structure-function studies on the bradykinin potentiating peptide
from Chinese snake venom (Agkistrodon halys pallas).";
Peptides 6 Suppl. 3:339-342(1985).
Peptides 6 Suppl. 3:339-342(1985).
PEPTIDE BOTH INHIBITS THE ACTIVITY OF THE
ANGIOTENSIN-CONVERTING ENZYME AND ENHANCES THAT INACTIVATE IT.
PRADYKININ BY INHIBITING THE KINASES THAT INACTIVATE IT.
ACTS AS AN INDIRECT HYPOTENSIVE AGENT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              pallas).
Eukaryota;
                                                                                                                                         "Synthesis
                                                                                                                                                                                                                                                                                                                                                                                                                            "Conserved amino acid sequence of a from coelenterates to humans.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Morphogenetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HUMAN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Lepidosauria; Squamata; Scleroglossa; Serpentes; Colubroidea; Viperidae; Crotalinae; Gloydius.
NCBI_TaxID=8714;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               fomo sapiens (Human)
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MOD RES 1 1
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    activator
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                                                                                                               Zachmann B., Bodenmuller is of a new neuropeptide, t. 131:317-321(1981).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GPPI
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(Rel. 34, Last
c neuropeptide
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    Druffel-Augustin
r acts as an autoc
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                                                                                                                                                                                                                                                  Sci. U.S.A.
                                                                                                                                                                              PubMed=7297679;
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                                                                                                                                                                                                                                                                                       sequence of a morphogenetic
    autocrine
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annotation updat
(Head activator)
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Pred.
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                       Dubel
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Dubel S. growth
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head acti
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    factor
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UH11 RAT
ID UH11 RAT
AC P56576;
DT 15-DEC-1998;
DT 15-DEC-1998;
DT 15-DEC-1998;
DT 15-DEC-1998;
DT 15-DEC-1998;
DE Unknown prote
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Best Local S
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Best Local S
Matches 2
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PIR; A01427; YHRT.
PIR; C01427; YHBO.
PIR; A93900; YHXAB.
PIR; B93900; YHJFHY.
                                                                                                                                                                                                                                                                                                           Spittaels K., Vankeerber Grauwels L., van Leuven
                                                                                                                                                                                                                 SEQUENCE
                                                                                                                                                                                                                                                                     Leptinotarsa decemlineata.";
Peptides 16:365-374(1995).
                                                                                                                                                                                                                                                                                       "Identification, characterization, a novel myotropic neuropeptide in (
                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Arthropoda; Mandibulata; Pancrustacea; Insecta; Pterygota; Neoptera; Endopterygota; Coleoptera; Pcucujiformia; Phytophaga; Chrysomeloidea; Chrysomelidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                 01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequence update)
01-NOV-1995 (Rel. 32, Last annotation update)
01-NOV-1995 (Rel. 32, Last annotation update)
Myotropic neuropeptide 1 (Led-MNP-I).
Leptinotarsa decemlineata (Colorado potato beetle).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MNP1 LEI
P42984;
                                                                                                                                                                                                                            Neuropeptide;
MOD_RES
                                                                                                                                                                                                                                                                                                                                                                                        Chrysomelinae; Leptinotarsa.
                                                                                                                                                                                                                                                                                                                                    MEDLINE=95380343; PubMed=7651886;
                                                                                                                                                                                                                                                                                                                                                           SEQUENCE,
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MOD_RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      in the G2/mitosis transition."; EMBO J. 8:3311-3318(1989).
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                                                                                                                                                                                                                                                 OVIDUCT
                                                                                                                                                                                                                                                          FUNCTION:
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CAUTION: THIS DEPTIDE WAS FIRST ISOLATED FROM NERVE AND WAS CALLED HEAD ACTIVATOR BY THE AUTHORS, BECAU HEAD-SPECIFIC GROWTH AND DIFFERENTIATION IN THIS AN BEEN FOUND IN MAMMALIAN INTESTINE AND HYPOTHALAMUS.
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             protein
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2; Conserv
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                         (Rel.
(Rel.
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Conservative
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van Leuven F., c
                                                                                                                                                                                                                             Amidation.
                                                                                                                                                                                                                                                          MYOTROPIC
                                                                     STANDARD;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cell cycle; Mitosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STANDARD;
37, Created)
37, Last sequence update)
37, Last annotation update)
5rom 2D-page of heart tissue
                                                                                                                                                                                                                 705 MW;
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                                                                                                                                                                                           22.8%;
                                                                                                                                                                                                                                                         PEPTIDE.
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de Loof A.;
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Pred. No.
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Pred. No.
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                                                                     PRT;
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the Colorado potato
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hes 0;
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THIS ANIMAL.
                                                                                                                                                                    Indels
             P11)
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            (Fragment).
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RESULT 15
AL16_CARMA
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Best Local S
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Jungblut P.R.;
Submitted (SEP-
                                                                          AL16 CARMA
P81819;
30-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AL15_CARMA STANDARD; PK1; G. FR. P81818; G. Created)
30-MAY-2000 (Rel. 39, Created)
30-MAY-2000 (Rel. 39, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
Carcinustatin 15.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rattus norvegicus (Rat).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rat
                                                                                                                                                                                                                                                                      -!- FUNCTION: MAY ACT AS A NEUROTRANSMITTER OR NEUROMODULATOR
-!- SIMILARITY: BELONGS TO THE ALLATOSTATIN FAMILY.

Neuropeptide; Amidation; Multigene family.

MOD RES

8

AMIDATION.

SEQÜENCE 8 AA; 811 MW; 922879D5AB47687D CRC64;
                                                                                                                                                                                                                                                                                                                                                                                      TISSUE=Cerebral ganglion, and Thoracic ganglion, MEDLINE=98121193; PubMed=9461295; Duve H., Johnsen A.H., Maestro J.-L., Scott A.G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Carcinus maenas (Common shore crab) (Green crab).

Eukaryota; Metazoa; Arthropoda; Mandibulata; Pancrustacea; Crustacea;
Malacostraca; Eumalacostraca; Eucarida; Decapoda; Pleocyemata;
Brachyura; Eubrachyura; Portunoidea; Portunidae; Carcinus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CARMA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRAIN-Wistar; TISSUE-Heart;
Li X.-P., Pleissner K.-P., Scheler C.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=10116;
                                                                                                                                                                                                                                                                                                                                      allatostatin superfamily in the shore crab
Eur. J. Biochem. 250:727-734(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NON
            Carcinus maenas (Common shore crab)
Eukaryota; Metazoa; Arthropoda; Mano
                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     UNSURE
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                                       Carcinustatin 16.
                                                30-MAY-2000 (Rel.
30-MAY-2000 (Rel.
30-MAY-2000 (Rel.
                                                                                                                                                                                                                                                                                                                                                                             Thorpe A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=6759;
                                                                                                                                                                                                                                                                                                                                                              Isolation and identification of multiple neuropeptides of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TER
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LLANEOUS: ON THE 2D-GEL THE DETERMINED PI OF THIS UNKNOWN
IN IS: 8.5, ITS MW IS: 42 kDa.
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            rab) (Green crab).
Mandibulata; Pancrustaçea; Crustacea;
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Best Local :
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Eur. J. Biochem. 250:727-734 (1997).
-:- FUNCTION: MAY ACT AS A NEUROTRANSMITTER
-:- SIMILARITY: BELONGS TO THE ALLATOSTATIN
Neuropeptide; Amidation; Multigene family.
MOD RES
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NCBI_TaxID=6759;
[1]
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(c) 1993 - 2003 Compugen Ltd.
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ALIGNMENTS

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01-MAY-1997
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EMBL; AF202074; AAF20919.1; -
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa;
Mammalia; Eutheria;
Bovidae; Bovinae; Bo
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SEQUENCE
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MEDLINE=97202105; PubMed=9049632;
Steelman S., Moskow J.J., Muzynski K., North C., Druck T.,
Steelman S., Moskow J.J., Daar I.O., Buchberg A.M.;
Montgomery J.C., Huebner K., Daar I.O., Buchberg A.M.;
Montgomery J.C., Huebner K., Daar I.O., Buchberg A.M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Wilm's tumor protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bos taurus (Bovine).
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01-MAY-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Xenopus laevis (African clawed frog).
Eukaryota; Metazoa; Chordata; Craniata;
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                                                                                                                                                                                    35.1%;
60.0%;
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Pred.
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Pred. No. 6.7e
2; Mismatches
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RESULT
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Best Local S
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Q9GJV3;
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Q1-MAR-2001
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SEQUENCE
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Q28866;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Cetacea; Mysticeti;
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    Lagenorhynchus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Balaenopteridae; Megaptera.
NCBI_TaxID=9773;
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01-MAY-1999
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Speciation, Systematics and Conservation.";
Submitted (APR-1999) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Lagenorhynchus obscurus
Eukaryota; Metazoa; Choi
Mammalia; Eutheria; Ceta
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01-MAR-2001 (TrEMBLrel.
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S73467; AAD14118.1; -.
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Conservative 0;
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(Dusky dolphin)
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Hare M.P., Cipriano F., Palumbi S.R.,
"Slow Evolution of Genetic Monophyly
                                                         01-MAR-2001 (TrEMBLrel.
01-MAR-2001 (TrEMBLrel.
01-MAR-2001 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Q9GJV2;
01-MAR-2001 (TrEMBLrel. 16,
01-MAR-2001 (TrEMBLrel. 16,
01-MAR-2001 (TrEMBLrel. 16,
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Mammalia; Eutheria;
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Mammalia; Eutheria;
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Lagenorhynchus acutus (Atlantic white-sided dolphin)
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
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                                                                                                                                      Q9GJV1
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Cetartiodactyla; Cetacea; Odontoceti
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Eukaryota; N
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Pseudomonas fluorescens.
Bacteria; Proteobacteria; gamm
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EMBL; AF140823; AAF98676.1; -.
EMBL; AF140824; AAF98677.1; -.
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                                 Metallothionein
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                                                                                                                                                                                                                                        Submitted (DEC-2001) to the
                                                                                                                                                                                                                                                       TRANSPOSON-TN5041A1, TN5041B, AND TN5041D1;
Kholodii G.Y., Gorlenko Z.M., Mindlin S.Z.,
"Distribution of distinct microvariants of
                                                                                                                                                                                                                                                                                   STRAIN=NC16-2,
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                                                                                                                                                                                                                                                                                                                                                                                 Q8RJF1
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NCBI_TaxID=90246;
                                                                                                                                                                                                                                                 bacteria.";
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AJ422129;
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Evolution of Genetic Monophyly in Dolphins: Impliation, Systematics and Conservation.";
tted (APR-1999) to the EMBL/GenBank/DDBJ databases
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                       STRAIN=JRS4;
Hogan D.A., Du P., Stevenson T.I.,
VanBogelen R.A.;
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MEDLINE=98083063; PubMed=9422600;
Martin K., Morlin G., Smith A., N
                                                                                                                                     Streptococcaceae;
                                                                                                                                                   Bacteria; Firmicutes; Bacillus/Clostridium
                                                                                                                                                                        Streptococcus pyogenes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        evidence for horizontal
                                                                                SEQUENCE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINB=97264487; PubMed=9110313; Cserjesi P., Fang H., Brandhorst "Metallothionein gene expression Lytechinus pictus."; Cytechinus pictus."; Med. Reprod. Dev. 47:39-46(1997).
                                                                                                                 NCBI_TaxID=1314;
                                                                                                                                                                                                                                                                      P82568;
                                                                                                                                                                                                                                                                                          P82568
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Martin K., Morlin G., Smith A., Nordyke A., Eisenstark A., "The tryptophanase gene cluster of Haemophilus influenzae evidence for horizontal gene transfer.";

J. Bacteriol 180:107-118(1998).
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01-DEC-2001
     Two-dimensional
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=7653;
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                                                                            AND MASS SPECTROMETRY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Proteobacteria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 9 AA; 1152 MW;
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(TrEMBLrel.
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   gel
                                                                                                                                     Streptococcus.
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13,
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Last annotation 
e (Fragment).
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Pred. No. 6.7e
1; Mismatches
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Last annotation updat
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                                     Whitton M.,
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No.
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2;
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                                                                                                                                                 group; Lactobacillales;
Streptococcus
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                                   Kilby G.W.,
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RESULT 14
Q93LX4
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Best Local S
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Matches 3
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                                                                                                                                                                                                                                                                                                                                                                                            Q93LX4;
Q93LX4;
01-DEC-2001
                                                                                                Submitted (MAY-2001) to the EMBL; AY034431; AAK61370.1; NON_TER 1
                                                                                                                               "Saha A., Haralalka S., Bhadra R.K.;
"Cloning and characterization of the origin of chromosome 1 of Vibrio cholerae.";
Submitted (MAY-2001) to the EMBL/GenBank/DDBJ of the
                                                                                                                                                                                                                                                                                                                                                           01-DEC-2001
01-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.

More M.I., Kirsch F.P., Rathjen F.G.;

"Targeted ablation of NrCAM and ankyrin-B resufibres leading to cataract formation.";

Submitted (FEB-2001) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Q99MG3;
Q99MG3;
01-JUN-2001
01-JUN-2001
                                                                               SEQUENCE
                                                                                                                                                                                                          SEQUENCE FROM N.A. STRAIN=0395;
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NON_TER
SEQUENCE
                                                                                                                                                                                                                                                                                  Bacteria; Proteobacteria;
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Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NrCAM protein
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Submitted (
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                                                                                                                                                                                                                                                                                                                                      (Fragment).
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   Similarity
3; Conserv
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                                                                               A
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                                                                             1303 MW;
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                   29.8%;
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50.0%;
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75.0%;
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MW=22592.04; MET
                                                                                                                                                                                                                                                                                  gamma
Score 17; DB; Pred. No. 9.1e
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Last
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Last annotation updat
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                                                                           464F014042D40337 CRC64;
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Pred. No.
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                                                                                                                                                                                                                                                                                subdivision; Vibrionaceae; Vibrio
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METHOD=ELECTROSPRAY.
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                 DB 2;
9.1e+03;
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Search completed: June 27, 2003, 17:33:04 Job time: 80 secs
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                                                                                                                         Query Match 29.8 Best Local Similarity 75.0 Matches 3; Conservative
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P82222;
P01-OCT-2001 (TrEMBLrel. 18, Created)
01-OCT-2001 (TrEMBLrel. 18, Last sequence update)
01-OCT-2001 (TrEMBLrel. 18, Last annotation update)
Unknown protein from 2D-page (Fragment).
Bombyx mori (Silk moth).
                                                                                                                                                                                                I Chuan Hsueh Pao 28:217-224(2001).

NON TER 10 10

SEQUENCE 10 AA; 1131 MW; 62CD78
                                                                                                                                                                                                                                                               SEQUENCE.
STRAIN=XINHANG X KEMING; TISSUE=BODY WALL, AND FAT BODY;
MEDLINE=21177481; PubMed=11280994;
Zhong B.X.;
Protein database for several tissues derived from five instar of
                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;
Pterygota; Neoptera; Endopterygota; Lepidoptera; Glossata; Ditrysia;
Bombycoidea; Bombycidae; Bombyx.
NCBI_TaxID=7091;
                                                                                                                                                                                                                                                     silkworm.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3 RVVER 7
||:|
5 RVINR 9
                                                                                              1 GPRV 4
                                                                    1 GPEV 4
                                                                                                                                                                                                10 AA; 1131 MW; 62CD783729D1B2CB CRC64;
                                                                                                                               29.8%; Score 17; DB 5; Length 10; 75.0%; Pred. No. 9.1e+03; tive 0; Mismatches 1; Indels
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                                                                                                                                Gaps
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Result
No.
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Maximum Match
Listing first
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Maximum DB
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Sequence:
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                                                                                                                                              Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution
      Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          seq length: 0
seq length: 11
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57
Match Length
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Query
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| SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1980.DAT:*
| SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1981.DAT:*
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/SIDS2/gcgdata/geneseq/geneseqp-embl/AA2002.DAT:*
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21.243 Million cell updates/sec
Description
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AAB15868	AAM49611	AAM99128	ABP18549	ABP12326	AAB83144	AAM99127	ABP20082	ABP12166	AAB67361	ABP22541	ABP20219	ABP14283	AAU95433	AAU95352	AAU94769	AAU94589	AAB27523	AAU95033	AAU94915	AAU94685	AAW49396	AAU95341	AAR96186	ABG35045	AAR96187	AAR96188	AAR96146	AAR96189	ABP18613	ABP12325	ABP20224	ABP12165	AAR96173	AAE02382
Human chemokine de	Human beta-defensi	relate	super	super n	ASB6411	related	motif	edns	Q.	HIV All motif env	motif	HIV A03 super moti	Human novel protei	novel	novel prote	or o	Ara h 1 linear IgE	Human novel protei	Human novel protei	٠.,	Human leucocyte an	Human novel protei		ndostatin	epi	n epi	ep.	rinogen epitop	B62 super	A02 super m	A03 motif	ñ	nogen epi	Human Factor VIII:

ALIGNMENTS

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RESULT 1
ABB76978
01-JUL-1999; 99FR-0008470.
                                       WPI; 2001-114394/13.
                                                       Serre G,
                                                                                                        01-JUL-1999;
                                                                                                                         05-JAN-2001.
                                                                                                                                         FR2795735-A1.
                                                                                                                                                         Homo sapiens.
                                                                                                                                                                         human; fibrinogen alpha chain.
                                                                                                                                                                                 Anti-arthritic; anti-inflammatory; fibrin; rheumatoid polyarthritis;
                                                                                                                                                                                                 Residues 36-46 of human fibrinogen alpha chain precursor.
                                                                                                                                                                                                                  22-JUL-2002
                                                                                                                                                                                                                                   ABB76978;
                                                                                                                                                                                                                                                  ABB76978 standard; peptide; 11 AA.
                                                                      (UYTO-) UNIV TOULOUSE SABATIER PAUL.
                                                      Sebbag M;
                                                                                                                                                                                                                 (first entry)
                                                                                                        99FR-0008470
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29.5 29

63.2 63.2 54.4 54.4 50.9

AAR65794 AAB01536 AAU85665 AAW92903

AAW65536 AAP71315 AAR06526

Residues 36-46 of Fibrinogen B-beta-Biotinyl-fibrin-an Sequence of fibrin reptide antigenic Fibrin-specific epplasmin substrate Sequence of

Plasmin substrate Fibrinogen ligand Factor VIII:C B do

New citrulline-containing polypeptide from fibrin, useful for diagnosis and treatment of rheumatoid polyarthritis $\,$

Example 1; Page 11; 23pp; French.

ABB76978 AAW11932

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RESULT 2
AAW11932
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                                                                                        A novel monoclonal antibody (MAb) reacts with human soluble fibrin, but not with human fibrinogen. Specifically when the MAb reacts with urea-treated des-AABB fibrin the reaction is not inhibited by peptides corresponding to fibrinogen A-alpha-chain residues 17-26 (AAW11931), B-beta-chain residues 15-24 (AAW11932) or gamma-chain residues 312-324 (AAW11933). The MAb is useful in immunoassays for soluble fibrin in plasma samples (e.g. by sandwich immunoassay), particularly for the diagnosis of pathological conditions such as
                                                         Sequence
                                                                                                                                                                                                                                                                       Monoclonal antibody with fibrinogen, is
                                                                                                                                                                                                                                 Example 3;
                                                                                                                                                                                                                                                             specimens
                                                                                                                                                                                                                                                                                                                  WPI; 1995-206667/27
                                                                                                                                                                                                                                                                                                                                                                                                   02-NOV-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                 01-NOV-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-MAY-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Monoclonal; antibody; human; soluble; fibrin; fibrinogen; urea-treated; des-AABB; B-beta-chain; immunoassay; diagnosis; disseminated intravascular coagulation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to a citrulline (Cit) containing polypeptide derived from all or part of the alpha- or beta-chains of fibrin by substitution of at least one arginine residue by Cit. The Cit containing polypeptides can be used for in vitro diagnosis of rheumatoid polyarthritis (RP), by detecting disease-specific autoantibodies, and therapeutically for neutralising the RP-associated autoimmune response. The present sequence is a fragment (residues 36-46) of the human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO9512617-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Monoclonal;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Fibrinogen B-beta-chain residues 15-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAW11932 standard; peptide; 10 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            fibrinogen alpha chain precursor, which was used in an example from the
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                 Similarity
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                                                                                                                                                                                                                                                                                                                                             Ito Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                  intravascular coagulation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                94WO-JP01844
                                                                                                                                                                                                                                 15;
                                                                                                                                                                                                                                                                   reactive with soluble human fibrin - but not useful for fibrin immunoassay in plasma
                                                                                                                                                                                                                            32pp; Japanese.
                                                                                                                                                                                                                                                                                                                                             Kohno I,
              93.0%; Score 53; 100.0%; Pred. No.
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Pred. No. 0.00014;
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DB 16; 1
. 0.00079;
thes 0;
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                          Length 10;
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10;

Conservative

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Mismatches

Indels

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Gaps

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The invention relates to organic compounds of formula V-L-R; V = organic Group having binding affinity for an angiotensin II receptor site; L = a CC linker group or a bond; R = a group detectable in in vivo imaging of a CC human or animal body; provided that when V is angiotensin or a peptidic CC angiotensin derivative or analogue, then V-L-R is not a non-metal CC radionuclide substituted peptide and L-V is not simply a peptide with a CC chelating agent amide bonded to a side chain. Also claimed are: (1) 1231-CC manufacture of a contrast medium for use in diagnosis involving CC deministration of the contrast medium to an animate subject and CC generating an image of at least part of the subject; (3) a method of CC generating an image of an animate subject comprising administration of a contrast medium to an animate subject and CC generating an image of an animate subject comprising administration of a contrast agent comprising the organic compound and generating the contrast agent comprising the organic compound and detecting the untake of the acent by anoitoring the organic associated with angiotensin II comprising administration of the organic compound and detecting the untake of the acent by anoitoring II.
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AAW65536
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                                                                                                                                                                                                                                                                                                   New angiotensin II receptor binding cardiovascular diseases and disorders
                                                                                                                                                                                                                                                                       Example 10; Page 88; 99pp;
                                                                                                                                                                                                                                                                                                                                                  WPI; 1998-348067/30
                                                                                                                                                                                                                                                                                                                                                                             Cuthbertson A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        07-MAY-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                           COCK/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             indothelin-1; angiotensin II receptor; imaging; cardiovascular disease;
coarran; contrast medium; diagnosis; lanthanide; biotin;
                                                                                                                                                                                                                                                                                                                                                                                                           COCKBAIN J
and detecting the uptake of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GPRVVERHQS
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97GB-0000699.
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96GB-0022365.
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                                                                                                                                                                                                                                                                       English.
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                                                                                                                                                                                                                                                                                                                   compounds
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    useful for imaging
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RESULT 4
AAP71315
ID AAP7
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                                                                                  FTTXXXXX
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                                         Query Match
Best Local S
Matches 7
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Best Local S
Matches 8
                                                                                                   The MAbs are specific to fibrin without fibrinogen cross-reactivity. They have increased binding to in vitro and in vivo thrombi. The MAbs can be used in immunoassays for fibrin in the presence of fibrinogen or other proteins. They can be used as immunoaffinity ligands for the purification of fibrin.

(Updated on 03-OCT-2002 to add missing OS field.)
                                                                                                                                                                                                  Screening of fibrin-specific monoclonal antibodies - by conwith immobilised crosslinked fibrin clot and screening with detectable labelling step
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          receptors, the administration and detection being optionally (and preferably) effected repeatedly. The organic compounds are useful for imaging cardiovascular diseases and disorders. The present sequence represents biotinyl-fibrin-anti-polymerant peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                    Misc-difference
                                                                                Sequence
                                                                                                                                                                                                                                                                  Matsueda GR,
                                                                                                                                                                                                                                                                                                                       14-APR-1986;
                                                                                                                                                                                                                                                                                                                                                                22-OCT-1987.
                                                                                                                                                                                                                                                                                                                                                                                     WO8706263-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Fibrin-specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence of fibrin immunogen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           03-OCT-2002
19-JUN-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAP71315;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAP71315 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                            Disclosure; Page 7; 41pp; English
                                                                                                                                                                                                                                             WPI; 1987-306855/43.
                                                                                                                                                                                                                                                                                       (GENO-)
                                                                                                                                                                                                                                                                                                                                           14-APR-1987;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (MAbs).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             l Similarity
8; Conserv
<u>بــ</u>
             1 GPRVVER 7
                                         Similarity 7; Conserv
                                                                                                                                                                                                                                                                                      GEN HOSPITAL
GPRVVER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GPRPPERHQS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GPRVVERHOS
                                                                                   7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10 AA;
                                         Conservative
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A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (updated)
(first entry)
                                                                                                                                                                                                                                                                 Haber E;
                                                                                                                                                                                                                                                                                                                      86US-0851514
                                                                                                                                                                                                                                                                                                                                           87WO-US00862
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      monoclonal antibody; screening
                                                                                                                                                                                                                                                                                                                                                                                                         /label= Arg-OH
                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
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                                                                                                                                                                                                                                                                                      CORP.
                                                 63.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                7 AA.
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                                                 Score 36;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 41; DB 19
Pred. No. 0.18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      for the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     prepn. of monoclonal antibodies
                                    DB b;
3. 7.8e+05;
0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19;
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                                                           Length
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                                         Indels
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AAR65794
ID AAR6
XX
AC AAR6
XX
DT 26 ---
XX
DE Fib
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AAR06526
ID AAR0
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DT 11-C
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KW fibr
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                                                                                      RESULT
                                                                                                                                                                           Query Match
Best Local
                                                                                                                                                                  Matches
                                                                                                                                                                                                                               Peptide is epitopic to fibrin, allowing anti-fibrin Abs raised without cross-reactivity to fibrinogen. They are particularly useful in detection of fibrin and thrombi.
                                                                                                                                                                                                                                                                                                 Fibrin-specific monoclonal antibodies - lacking fibrinogen.cross-reactivity, obtd. using peptide(s) comprising fibrin-specific epitopic sequences.
                                           AAR65794;
                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                             Claim 1; Page 17; 12pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                23-APR-1984;
30-JAN-1986;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAR06526;
                       26-JUN-1995
                                                                AAR65794 standard;
                                                                                                                                                                                                                                                                                                                                               WPI; 1990-185723/24.
                                                                                                                                                                                                                                                                                                                                                                      Matsueda
                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-JAN-1986;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22-MAY-1990.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US4927916-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         fibrin; fibrinogen; thrombi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11-OCT-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAR06526 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                           (GEHO-)
                                                                                      σ
                                                                                                                                      1 GPRVVER: 7
                                                                                                                                                                                                                                                                                                                                                                                           GEN
                                                                                                                                                                 Similarity 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antigenic to
                                                                                                                                                                                                                                                                                                                                                                     GR,
                                                                                                                      GPRVVER 7
                                                                                                                                                                                                             7 AA;
                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                           HOSPITAL CORP
                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first
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                                                                                                                                                                                                                                                                                                                                                                                                                84US-0603155
86US-0824228
                                                                                                                                                                                                                                                                                                                                                                                                                                                86US-0824228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /label=Lys or Arg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      label=Asp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /label=Leu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /label=Pro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       label=His or Pro
                                                                peptide; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   entry)
                                                                                                                                                                            63.2%; Score 36; 100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         immunoaffinity.
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                                                                                                                                                                  Mismatches
                                                                                                                                                                           DB 11; I
7.8e+05;
                                                                                                                                                                                      Length
                                                                                                                                                                  Indels
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                                                                                                                                                                  Gaps
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Fibrin-specific epitopic peptide.

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RESULT 7
AAB01536
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Best Local
                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                 Polymer; biomaterial; conjugate; hydrogel; drug delivery; adhesive; sealant; tissue engineering; wound healing; scaffold; cell transplant; adhesion prevention; cell migration; collagenase;
(HUBB/) HUBBELL J A
                        01-FEB-1999;
                                          01-FEB-2000; 2000WO-US02608
                                                                 03-AUG-2000
                                                                                     WO200044808-A1
                                                                                                          Synthetic.
                                                                                                                               plasmin; elastase
                                                                                                                                                                               Plasmin substrate site found
                                                                                                                                                                                                    08-NOV-2000
                                                                                                                                                                                                                           AAB01536;
                                                                                                                                                                                                                                                                                                                                                                                                           AAR65789-R65794 are synthetic peptides comprising fibrin-specifically sequences, they can be used to prepare hybridoma cell lines, which produce antifibrin-specific monoclonal antibodies substantially devoid of fibrinogen cross-reactivity. These antibodies are useful in the in vivo and in vitro detection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-JAN-1986;
22-DEC-1989;
24-AUG-1992;
                                                                                                                                                                                                                                             AAB01536 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic epitopic peptide(s) of variable length - capable of eliciting fibrin specific antibodies free of fibrinogen
                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                     antibodies are useful in the in of thrombi and fibrin deposits.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 9; Column 14; 12pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cross-reactivity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1994-332411/41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Haber
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23-APR-1984;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23-APR-1984;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US5357042-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Fibrin-specific epitopic peptides; thrombi detection; fibrinogen; antifibrin-specific monoclonal antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18-OCT-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GEHO ) GEN HOSPITAL CORP
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                                                                                                                                                                                                                                                                                                 μ
                                                                                                                                                                                                                                                                                                                                         7; Conserv
                                                                                                                                                                                                                                                                                                                     GPRVVER 7
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                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                   ΑA,
                                                                                                                                                                                                   (first entry)
                        99US-0118093.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        84US-0603155.
86US-0824228.
89US-0454954.
92US-0932729.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            84US-0603155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matsueda GR
                                                                                                                                                                                                                                            peptide; 6
                                                                                                                                                                                                                                                                                                                                                   63.2%;
                                                                                                                                                                                                                                                                                                                                      Score 36; DB: Pred. No. 7.8
                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                             in fibrinogen.
                                                                                                                                                                                                                                                                                                                                      DB 15; I
. 7.8e+05;
ches 0;
                                                                                                                                                                                                                                                                                                                                                          Length 8;
                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                            fibrin-specific
ybridoma cell
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RESULT 8
AAU85665
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                     Biomaterial; vasotropic; anticoagulant; thrombolytic; vulnerary; infection; adhesion; thrombosis; restenosis; adhesive;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                transplant devices. The blomaterials are also useful for adhesion prevention to minimise unwanted operative or post-traumatic adhesions. In the instances where the precursor for the polymeric biomaterial is a peptide, the nucleophile is the amino acid cysteine. This results in peptides of structure similar to H_2N-CXXXXXCXXXXC-COOH (See GENESEQ records AABOIS31-B01535). The length of XXXXX is variable and can be of any number. It is particularly useful when the sequences in the domains shown as XXXXX are substrates for enzymes that are involved in cell migration (e.g. as substrates for enzymes such as collagenase, plasmin or elastase).
            (EIDG-) EIDGENOESSISCHE TECH HOCHSCHULE ZUERICH. (UYZU-) UNIV ZURICH.
                                                    02-JUN-2000; 2000US-0586937.
                                                                                 04-JUN-2001;
                                                                                                              06-DEC-2001
                                                                                                                                                                    Unspecified
                                                                                                                                                                                                               vulnerary; infe
sealant; tissue
                                                                                                                                                                                                                                                                       Plasmin
                                                                                                                                                                                                                                                                                                   21-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               self selective reaction between a strong nucleophile and a conjugated unsaturated bond or a conjugated unsaturated group, by nucleophilic addition. The polymeric hydrogels can be used in a variety of applications. They can be used to deliver therapeutic molecules to a subject, as adhesives or sealants (e.g. sealing air leaks on the lung), as tissue engineering and wound healing scaffolds, and as cell transplant devices. The biomaterials are also useful for adhesion
                                                                                                                                        WO200192584-A1
                                                                                                                                                                                                                                                                                                                                AAU85665;
                                                                                                                                                                                                                                                                                                                                                        AAU85665 standard; Peptide; 6 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         combining peptide) o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Producing polymeric biomaterials by polymerizing components (e.g. polymer, protein or peptide) of useful for delivering therapeutic molecules to a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hubbell
Tirelli
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 44; 119pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              adhesives or sealants
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2000-524289/47.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ymerisation of the two components. Polymerisation occurs through
                                                                                                                                                                                                transplant
                                                                                                                                                                                                                                                                       substrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         <u>ه</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ŊĄ,
                                                                                                                                                                                                                                                                                                                                                                                                                                 GPRVVE
                                                                                                                                                                                                                                                                                                                                                                                                                                                           GPRVVE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     g two or more precursor components (e.g. I of the biomaterial under conditions that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                of making polymeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               6 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                2001WO-US18101.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Vernon B;
                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Elbert D,
                                                                                                                                                                                                            engineering;
                                                                                                                                                                                                                                                                     site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   54.4%;
                                                                                                                                                                                                                                                                     found
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 31;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              English
                                                                                                                                                                                                               wound
                                                                                                                                                                                                                                                                     in fibrin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             biomaterials is described comprising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Σ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                          healing
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                                                                                                                                                                                                                                                                     #1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    7.8e+05;
thes 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Þ
                                                                                                                                                                                                            scaffold;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Schoenmakers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      two or more precursor the biomaterial, subject and as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               polymer, protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
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Hubbel JA,

Elbert D,

Schoenmakers

an ester

for medical treatment an ester or amide bond

comprises an active or a onto the active or binding

relates to a biomaterial comprising an active

or a binding

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RESULT 9
AAW92903
ID AAW9
XX AAW9
AC AAW9
XT 14-M
XT 14-M
XX Fibr
XX Fibr
XX Fibr
KW micr
KW hear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local S
Matches 6
                                                                                                                                                                                                                                                                                          disease, disorder or infection in a mammal e.g. human, for preventing adhesions, thrombosis or restenosis in a mammal and for delivering the active compound (preferably protein or peptide) to a cell, tissue, organ system or a body of a mammal as adhesives or sealants, as tissue engineering and wound healing scaffolds and as cell transplant devices. The bond of the biomaterial has a half life of 1 hour - 1 year (preferably 1 day - 9 months, especially 2 days - 2 months) in an aqueou solution at pH 7.4 and 37plusoc. Thus the biomaterial releases the therapeutic compound over a clinically relevant time-frame. The present sequence is a peptide which acts as a therapeutically active compound suitable for inclusion in the biomaterial of the
                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                          -swellable polymer containing conjugated unsaturated groups;
U = product of the addition of a nucleophile to an electrophile that
is attached to the water-soluble or water-swellable polymer; and
n = 2 - 3. The biomaterials are useful for treating or preventing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 81; 222pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New biomaterial useful binding group and has a
                                                                  14-MAY-1999
                                                                                                            AAW92903 standard; peptide; 9
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Fibrinogen; ligand; fibrin; treatment; prevention; rheology; diabetes; microcirculatory disorder; retinopathy; polyneuropathy; apoplexy; sepsis; hearing loss; sepsis, occlusive arterial disease; kidney function.

Fibrinogen ligand peptide #7.

(first

entry)

Similarity 6; Conserv

Conservative

54.4%; Score 31; DB 100.0%; Pred. No. 7. Live 0; Mismatches

DB 23, o. 7.8e+05; 0;

Length 6; Indels

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Best Local :
         03-DEC-1993;
07-FEB-1991;
16-MAY-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                        for preparation of an agent for treating or preventing microcirculatory disorders and/or for modifying rheology, in mammals. The products of the invention can be specifically used to treat or prevent disorders associated with diabetes, retinopathy, polyneuropathy, apoplexy, hearing loss, sepsis, occlusive arterial disease and/or impaired kidney function Reducing the amount of fibrinogen and/or fibrin in the blood alleviates
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Using ligand for fibrinogen or fibrin to treat or prevent microcirculatory disorders - and to control blood rheology, e.g. diabetes related conditions, also adsorber material with bound
                                                                                                                                                                           Factor VIII:C B domain C-terminal peptide.
                                                                                                                                                                                                   29-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 17; 23pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bieber
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     02-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                            23-AUG-2001
                                                                                                                     Homo sapiens
                                                                                                                                                      Factor VIII:C; human; haemophilia A; blood clotting; therapy;
                                                                                                                                                                                                                        AAB82729;
                                                                                                                                                                                                                                              AAB82729 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This invention describes the use of a ligand for fibrinogen and/or fibrin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1999-120788/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO9902565-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Modified-site
                                                                                                 US2001016340-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (THER-) THERASORB MEDIZINISCHE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    '11
                                                                                                                                                                                                                                                                                                                 1 GPRVVERHQS
                                                                                                                                                                                                                                                                                                                                                                                                                 symptoms
                                                                                                                                                                                                                                                                                                    GPR-PERHES
                                                                                                                                                                                                                                                                                                                                                                                           9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    the process
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Koll R,
                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                  (first entry)
                                                      2000US-0748062
93US-0161770.
91US-0652099.
95US-0441943.
86US-0822989.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               97DE-1029591
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    98WO-EP04090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note= "C-terminal amidated"
                                                                                                                                                                                                                                              Peptide;
                                                                                                                                                                                                                                                                                                                          10
                                                                                                                                                                                                                                                                                                    9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Richter W,
                                                                                                                                                                                                                                                                                                                                                          51.8%;
70.0%;
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                                                                                                                                                                                                                                                                                                                                                          Score 29.5;
Pred. No. 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SYSTEME GMBH
                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tschoepe W;
                                                                                                                                                                                                                                                                                                                                                          .8e+05
                                                                                                                                                                                                                                                                                                                                                                      BB
                                                                                                                                                                                                                                                                                                                                                                     20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                03-DEC-1993;
07-FEB-1991;
27-JAN-1986;
19-MAY-1987;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence is that of the C-terminal sequence of the B domain of human Factor VIII.C. The invention relates to recombinant protein complexes having Factor VIII.C activity, and methods for their production in eukaryotic host cells. The recombinant protein complexes comprise a first polypeptide homologous to human Factor VIII.C A domain and a second polypeptide homologous to human Factor VIII.C A domain and a second polypeptide homologous to human Factor VIII.C A domain sequence may be extended at its C-terminal end by a B domain sequence comprising the N-terminal sequence of human Factor VIII.C B domain (see AAB82727), a hinge peptide (see AAB82728) homologous to a human immunoglobulin heavy chain hinge region, and the B domain fortune of the protein complexes are considered.
                                                                                                                                                                                                                                           Human; Factor VIII:C; FVIII:C; coactype A haemophilia; blood clotting recombinant protein; immunogen.
(CHIR ) CHIRON CORP.
(NOVO ) NOVO NORDISK AS.
                                                                                                                                                       08-MAY-2001
                                                                                                                                                                                                                                                                                                      Human Factor VIII:C B-domain C-terminal sequence
                                                                                                                         16-MAY-1995;
                                                                                                                                                                                      US6228620-B1
                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                     10-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                             AAE02382 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    chromatography, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C-terminus (present sequence). The protein complexes are useful treating classical (Type A) haemophilia and other blood clotting disorders. They are also useful as immunogens for producing annibodies, for isolating von Willebrand factor by affinity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 12; Page 27; 29pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (CHIR )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New polypeptide having Factor VIII:C (Type A) haemophilia and other blood immunogens for producing antibodies, for Factor VIII:C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-535541/59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Chapman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19-MAY-1987;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CHIRON CORP.
NOVO NORDISK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRVVERHQ 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PPVLKRHO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Burke RL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                   (first entry)
                                              93US-0161770.
91US-0652099.
86US-0822989.
87US-0051916.
                                                                                                                         95US-0441943
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       87US-0051916
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  in diagnostic assays for Factor VIII:C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              50.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rasmussen ME,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           English.
                                                                                                                                                                                                                                                                                                                                                                                                9 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 29; DB 22;
Pred. No. 7.8e+05;
                                                                                                                                                                                                                                                     coagulation; von-Wille-brand factor;
ting disorder; haemostatic; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 activity, for treating classical clotting disorders, for use as or for use in diagnostic assays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mikkelson JM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
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RESULT 12
AAR96173
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Best Local S
Matches 5
WPI; 1996-251888/25
                            Boger JS, Dahlgren ME,
                                                                                            06-JUN-1995;
07-NOV-1994;
                                                                                                                                           03-NOV-1995;
                                                                                                                                                                                                        WO9614580-A1
                                                                                                                                                                                                                                                                                                                             Epitope; cleavage product; human; leukocyte elastase; HLE; fibrinogen; monoclonal antibody; probe; detection; antigen; blood; peritoneal flui sputum; broncheoalveolar lavage fluid; assay; inhibitor; arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAR96173 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             encoded by the polynucleotide present in plasmid pSVF8-200. The composition is useful as an immunogen for the production of antibodies, for isolating von-Wille-brand factor by affinity chromatography, in diagnostic assays for Factor VIII:C and for treative present sequence is C-terminal fragment of human Factor VIII:C B-domain used in the recombinant protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The patent relates to a nucleic acid composition for introducing nucleic acid into a eukaryotic host cell to obtain expression of a recombinant protein which lacks all or a portion of the B domain or human Factor VIII:C (hFVIII:C) and is capable of coagulation active the recombinant protein consists of a polypeptide substantially homologous to mature A domain of hFVIII:C as encoded by the polynucleotide present in plasmid pSVF8-200 (ATCC) number 40190); a second polypeptide homologous to mature C domain of hFVIII:C as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New nucleic acid complexes having factor VIII activity and coagulation activity, for treating hemophilia and other blood clotting disorders, or as immunogens for the production of antibodies -
                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                   chronic obstructive pulmonary disease; myelogenous leukaemia;
                                                                                                                                                                                                                                                                                                                                                                                                 Fibrinogen
                                                                                                                                                                                                                                                                                                                                                                                                                                 19-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAR96173;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                   pulmonary emphysema; chronic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-335014/35
                                                                                                                                                                                                                                                                                      nfantile respiratory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Chapman B,
                                                                                                                                                                                                                                                                   respiratory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     <u>ن</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PPVLKRHQ 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRVVERHQ 9
                                                                                                                                                                                                                                                                                                                                                                                            epitope, represents alpha chain residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Column 51-52; 33pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Burke RL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                           95US-0469141
94US-0335524
                                                                                                                                           95WO-US13794
                                                             გ
                                                                                                                                                                                                                                                                     distress
                                                             INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   50.9%;
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                             Davies DTP,
                                                                                                                                                                                                                                                                     syndrome
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 29;
Pred. No.
                                                                                                                                                                                                                                                                                                                age fluid; assay; inhibitor; arthritis;
bronchitis; cystic fibrosis; bronchiectasis;
                                                                                                                                                                                                                                                                                  syndrome; gout;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; DB 22; LC
5. 7.8e+05;
1;
                             Humes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mikkelson
                          ŢĽ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 9;
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                             Mumford
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                            ß
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            for treating
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         activity.
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RESULT 13
ABP12165
ID ABP12
XX ABP12
XX ABP12
XX ABP12
XX HIV A
XX HIV A
XX HIV A
XX HIV A
XX HOUN
CW ANTIG
CO The
PN WO200
XX CEPIN
XX WPI;
PT Vacc:
PT Pept.
XX Clail
XX Clail
XX Clail
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The sequences given in AAR96164-81 represent antigenic peptides derived from the N-terminus of the primary cleavage products of human leukocyte elastase (HLE) cleaved human fibrinogen. These epitopes are capable of inducing specific antibodies and acting as specific probes for the detection of antibodies. Monoclonal antibodies reactive against these peptides, and other HLE cleaved fibrinogen derived peptides (see also AAR96146-63 and AAR96182-94) are prepared by standard techniques. These monospecific antibodies may be used to assay for the formation of complimentary cleavage product antigens or epitopes in whole blood or other body fluids, peritoneal fluid, sputum or broncheoalveolar lavage fluid. The assay for cleavage products is dependent upon the presence of HLE in the sample. This assay can also be used for the evaluation of HLE inhibitors. The antibodies may be used to diagnose
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New isolated fibrinogen derived cleavage products - used for detection of leukocyte elastase activity in disease diagnosis and for evaluating elastase inhibitors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          and monitor diseases such as arthritis, gout, pulmonary emphysema, chronic bronchitis, cystic fibrosis, chronic obstructive pulmonary disease, bronchiectasis, adult or infantile respiratory distress
                                                                                                                                                                                                                                                                                                                                                                                  Human immunodeficiency virus type 1.
                                                                                                                                                                                                                                                                                                                                                                                                                  antigen; vaccine; HIV infection; immunisation; virucide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HIV A02 super motif env peptide #222
                                                                                                                                WPI; 2001-354887/37
                                                                                                                                                                                  Sette
                                                                                                                                                                                                                                                  05-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABP12165
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABP12165 standard; Peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           syndrome and myelogenous leukaemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 21; 109pp; English.
                                              Claim 32; Page 118; 448pp; English
                                                                                 peptide
                                                                                                                                                                  Baker
                                                                                                                                                                                                                                                                                 05-OCT-2000; 2000WO-US27766
                                                                                                                                                                                                                                                                                                                    12-APR-2001
                                                                                                                                                                                                                                                                                                                                                     WO200124810-A1
                                                                                                                                                                                                                   (EPIM-)
                                                                                                                                                                                                                                                                                                                                                                                                                                 HIV-1; human immunodeficiency virus; env; pol; vif; tat; cytotoxic T lymphocyte; CTL; immune;
                                                                                                                                                                 D.A
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5; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       6
                                                                               groups, useful for vaccinating
                                                                                                                                                                                                                EPIMMUNE INC
                                                                                                                                                             Sidney J,
Celis E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ERHQS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ERHQS 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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                                                                                                                                                                 Southwood Kubo RT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%;
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                                                                                                                                                             S, L
                                                                               human immunodeficiency inating against HIV-1 -
                                                                                                                                                                 Livingston ey HM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 17; 1
7.8e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         respiratory distress
                                                                                                                                                                               BD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                   response; epitope;
                                                                                                                                                                               Chesnut
                                                                                                 virus-1 (HIV-1)
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The present invention describes a composition (I) comprising a prepared human immunodeficiency virus-1 (HIV-1) group comprising an amino acid

against HIV-1

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RESULT 14
ABP20224
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SXFXGXGXA
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Best Local
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Baker DM,
          The present invention describes a composition (1) comprising fundan immunodeficiency virus-1 (HIV-1) group comprising an an sequence selected from 51 defined amino acid sequences (ABL25 ABP25397). (1) has virucide activity and can be used in vacci
                                                                                                                          Vaccine compositions comprising human immunodeficiency virus-1 (HIV-1) peptide groups, useful for vaccinating against HIV-1 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HIV;
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                                                                                               Claim
                                                                                                                                                                              WPI; 2001-354887/37.
                                                                                                                                                                                                                                                                                                 05-OCT-1999;
                                                                                                                                                                                                                                                                                                                                  05-OCT-2000; 2000WO-US27766
                                                                                                                                                                                                                                                                                                                                                                   12-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                    Human immunodeficiency virus type
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HIV A03 motif env peptide #428
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antigen; vaccine;
                                                                                                                                                                                                                                                                (EPIM-) EPIMMUNE INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HIV-1; human immunodeficiency virus; env; pol; gag; nef; vpr; vif; tat; cytotoxic T lymphocyte; CTL; immune response; epito
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Celis E,
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Kubo RT, Grey
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66.78;
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Pred. No.
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group-based vaccines has several advantages over traditional vaccines, particularly when compared to the use of whole antigens in vaccine

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   human immunodeficiency virus-1 (HIV-1) group comprising an amino acid sequence selected from 51 defined amino acid sequences (ABL25347 to ABP25397). (I) has virucide activity and can be used in vaccines. (I) may be used for immunising subjects against HIV-1 infections. The use of group-based vaccines has several advantages over traditional vaccines, particularly when compared to the use of whole antigens in vaccine
                                                                                                                                                                                                                                                                            Sette A,
Baker DM,
                                                                                                                                                                               Vaccine compositions comprising human immunodeficiency virus-1 (HIV-1) peptide groups, useful for vaccinating against HIV-1 -
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                                                                                                          The present invention describes a composition (I) comprising a prepared
                                                                                                                                                  Claim 32; Page 121; 448pp; English.
                                                                                                                                                                                                                                         WPI; 2001-354887/37
                                                                                                                                                                                                                                                                                                                                                                                                05-OCT-2000; 2000WO-US27766.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HIV; HIV-1; human immunodeficiency virus; env; pol; gag; nef; vpr; vpu; vif; tat; cytotoxic T lymphocyte; CTL; immune response; epitope; antigen; vaccine; HIV infection; immunisation; virucide.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       represent peptide sequences used in the exemplification of the present
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Job time :

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Search completed: June 27, 2003, 17:31:38

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Best Local S
Matches 6
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                                                                                                                                                 Sequence
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Post-processing: Minimum Match
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Maximum DB
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Perfect score:
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    protein search, using sw model
 is the number of results predicted by chance to have a ter than or equal to the score of the result being printed,
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/cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*
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/cgn2_6/ptodata/1/pubpaa/US07_NEW_PUB.pep:*
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0 US-09-725-715-7
0 US-09-748-062-34
US-10-062-710-91
US-10-208-806-63
US-10-228-806-63
US-09-954-385-26
US-10-228-806-62
US-09-955-548A-180
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US-09-955-548A-180
US-09-955-958-23
US-10-066-474-23
US-10-066-474-23
US-09-951-809C-9
0 US-09-9845-849-9
US-10-058-513-26
US-09-903-185-4
US-10-185-425-21
US-09-876-904A-188
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           GENERAL INFORMA
Sequence 10, Appl
Sequence 7, Appl
Sequence 34, Appl
Sequence 28, Appl
Sequence 28, Appl
Sequence 63, Appl
Sequence 63, Appl
Sequence 62, Appl
Sequence 180, Appl
Sequence 23, Appl
Sequence 23, Appl
Sequence 278, Appl
Sequence 4, Appli
Sequence 4, Appli
Sequence 4, Appli
Sequence 4, Appli
Sequence 21, Appli
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 Sequence
 188,
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US-09-953-657-5
GENERAL INFORMATION:
                                              ISSUE:
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	136,	Sequence 60, Appl	193,	'n	60,	48,	Sequence 47, Appl	Sequence 525, App		12,	54		•	603,	Sequence 596, App	482,	69,	661,	547,	451,	350,	161,	58,	Sequence 28, Appl	Sequence 49, Appl

ALIGNMENTS

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APPLICATION NUMBER: 08/595,718
FILING DATE: «Unknown»
ATTORNEY/AGENT INFORMATION:
NAME: Giblin, James A.
REGISTRATION NUMBER: 25772
REGISTRATION NUMBER: MSB-723:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (510)705-7910
                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/09/953,657
FILING DATE: 17-Sep-2001
CLASSIFICATION: «Unknown»
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM
OPERATING SYSTEM: DOS
SOFTWARE: WORDERFECT 6.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Baumbach, George A.
Hammond, David J.
TITLE OF INVENTION: Fibrinogen Binding Peptides
NUMBER OF SEQUENCES: 8
SEQUENCE DESCRIPTION: SEQ ID NO: 5:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.44Mb Storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: Bayer Corporation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Buettner, Joseph A.
Dadd, Christopher A.
                                                                                                    TELEFAX: (510)705-7904
JOURNAL: Thromb. Hae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STATE: California COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Berkeley
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 94701-1986
                    28-JUN-1990
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US-09-925-715-7
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US-09-765-614B-10
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                                                                                                                                                       FILE REFERENCE: REF/Klaveness/206
CURRENT APPLICATION NUMBER: US/09/925,715
CURRENT FILING DATE: 2001-08-10
NUMBER OF SEQ ID NOS: 27
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 7
                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Nycomed Imaging AS
TITLE OF INVENTION: Improvements in
TITLE OF INVENTION: agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILE REFERENCE: REF/Klaveness/054
CURRENT APPLICATION NUMBER: US/09/765,614B
CURRENT FILING DATE: 2001-07-10
NUMBER OF SEQ ID NOS: 31
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 10
LENGTH: 10
TYPE: PRT
                                                                                                                                                                                                                                                                                                                      Sequence 7, Application US/09925715
Patent No. US20020102217A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: Description of Art OTHER INFORMATION: Sequence: Biotinyla OTHER INFORMATION: fibrin-anti-polyme NAME/KEY: MOD RES LOCATION: (1)
OTHER INFORMATION: Biotinylated-Gly NAME/KEY: MOD RES LOCATION: (10)
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Best Local
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Best Local Similarity
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APPLICANT: Nycomed
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                           OTHER INFORMATION: Description of Artificial Sequence:Biotinylated OTHER INFORMATION: fibrin-antipolymerant peptide NAMEKEY: MOD_RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Nycomed Imaging AS
TITLE OF INVENTION: Improvements in or relating
TITLE OF INVENTION: diagnostic/therapeutic
TITLE OF INVENTION: agents
                                                                                                  ORGANISM: Artificial Sequence
                                                                                                                          TYPE: PRT
OTHER INFORMATION: Biotin-Gly
                                                                                           FEATURE:
                                                                                                                                           ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: AMIDATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Description of Artificial Sequence:Biotinylated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 fibrin-anti-polymerant peptide
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88.9%;
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Pred. No. 3
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Pred. No. 0
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US-09-748-062-34
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OTHER INFORMATION: AMIDATION US-09-925-715-7
                                   Query Match
Best Local Similarity
Matches 5; Conserv
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Best Local Similarity
Matches 8; Conserv
                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO: 34:
                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/441,943
FILING DATE: <Unknown>
APPLICATION NUMBER: US 822,989
FILING DATE: 27-JAN-1986
ATTORNEY/AGENT INFORMATION:
NAME: BAROUSKY, KENNETH
                                                                                                               TOPOLOGY: linear
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 34:
                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                  REGISTRATION NUMBER: 36,442
REFERENCE/DOCKET NUMBER: 2300-0048.10
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0 CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: PROTEIN COMPLEXES HAVING FACTOR VIII:C ACTIVITY AND PRODUCTION THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
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ADDRESSEE: REED &
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PRVVERHO 9
                                                                                                                                                                                 TYPE: amino acids
                                                                                                                                                                 STRANDEDNESS: single
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FILING DATE: 22-Dec-2000
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                    TELEPHONE: (415) 327-3400
TELEFAX: (415) 327-3231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: PALO ALTO
STATE: CALIFORNIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: F1
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                                 Conservative
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RASMUSSEN, MIRELLA EZBAN
MIKKELSON, JAN MOLLER
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                                               50.9%;
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                               Score 29; DB 10;
Pred. No. 3.8e+05;
2; Mismatches 1
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Pred. No.
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                                                                                                                                                                     RESULT 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn version 3.1
SEQ ID NO 28
LENGTH: 10
                                                                                 Sequence 63, Application US/10228806
Publication No. US20030049237A1
GENERAL INFORMATION:
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Best Local Similarity
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Publication No. US20030108554A1
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Best Local
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APPLICANT: Chu, Yong-Liang
APPLICANT: Chu, Yong-Liang
APPLICANT: Qiu, Jian-Tai
TITLE OF INVENTION: Polymeric Conjugates for Delivery of
TITLE OF INVENTION: MHC-Recognized Epitopes
TITLE OF INVENTION: Via Peptide Vaccines
FILE REFERENCE: 3781-001-27
CURRENT APPLICATION NUMBER: US/10/062,710
CURRENT FILING DATE: 2002-02-05
APPLICANT: Bannon, et al.
TITLE OF INVENTION: Methods and Reagents for Decreasing Clinical Reactions
TITLE OF INVENTION: to Allergy
FILE REFERENCE: 2002834-0043
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Revert-Ros, Francisco
TITLE OF INVENTION: GIPs, a Family of Polypeptides with Transcription Factor Activity
TITLE OF INVENTION: Interact with Goodpasture Antigen Binding Protein
FILE REFERENCE: 98,723-F-US
CURRENT APPLICATION NUMBER: US/10/309,851
CURRENT FILING DATE: 2002-12-04
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PRIOR FILING DATE: 2001-08-08
NUMBER OF SEQ ID NOS: 232
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TYPE: PRT
ORGANISM: Artificial Sequence
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Pred. No.
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Pred. No. le+02;
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RESULT 9
US-10-228-806-62
; Sequence 62, Application US/10228806
; Publication No. US20030049237A1
; GENERAL INFORMATION:
; APPLICANT: Bannon, et al.
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; ORGANISM: Arachis hypogaea
US-10-228-806-62
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                                                             TITLE OF INVENTION: Methods and Reagents for Decreasing Clinical Reactions TITLE OF INVENTION: to Allergy FILE REFERENCE: 2002849-0043 CURRENT APPLICATION NUMBER: US/10/228,806 CURRENT FILING DATE: 2002-08-26 NUMBER OF SEQ ID NOS: 81 SOFTWARE: Patentin Ver. 2.1 SEQ ID NO 62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Aehle, Wo
APPLICANT: Baldwin,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 26
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LENGTH: 10
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CURRENT FILING DATE: 2002-08-26
NUMBER OF SEQ ID NOS: 81
SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILE REFERENCE: GC690
CURRENT APPLICATION NUMBER: US/09/954,385
CURRENT FILING DATE: 2001-09-12
NUMBER OF SEQ ID NOS: 433
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Winetzky, Deborah S.
TITLE OF INVENTION: Binding Phenol Oxidizing Enzyme-peptide
TITLE OF INVENTION: Complexes
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                                 TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
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                                               LENGTH: 10
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                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
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Van Gastel, Franciscus J.C.
Janssen, Giselle G.
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Murray, Christopher J.
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                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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55.6%;
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Pred. No.
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APPLICANT: Nicolette, Charles A.
TITLE OF INVENTION: THERAPEUTIC ANTI-MELANOMA COMPOUNDS
FILE REFERENCE: GZ 2103.00
CURRENT APPLICATION NUMBER: US/09/922,405B
CURRENT ELING DATE: 2002-06-10
PRIOR APPLICATION NUMBER: US 60/23,641
PRIOR FILING DATE: 2000-08-04
PRIOR PRIOR APPLICATION NUMBER: US 60/255,502
PRIOR FILING DATE: 2000-12-13
PRIOR APPLICATION NUMBER: US 60/264,432
PRIOR APPLICATION NUMBER: US 60/279,005
PRIOR FILING DATE: 2001-01-25
PRIOR APPLICATION NUMBER: US 60/279,005
                                                         ; ORGANISM: Homo sapien
US-09-922-405B-23
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Query Match
Best Local Similarity
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Sequence 23, Application US/09922405B
Publication No. US20020182218A1
GENERAL INFORMATION:
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                                                                                                                             SEQ ID NO 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/865,548A CURRENT FILING DATE: 2001-05-16 PRIOR APPLICATION NUMBER: US 60/290,958 PRIOR FILING DATE: 2001-05-16 NUMBER OF SEQ ID NOS: 204 SOFTWARE: Patentin version 3.1 SEQ ID NO 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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Best Local Similarity
Matches 5; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Admon, Arie
TITLE OF INVENTION: METHOD OF IDENTIFYING PEPTIDES CAPABLE OF BINDING TO MHC MOLECUL
TITLE OF INVENTION: PEPTIDES IDENTIFIED THEREBY AND THEIR USES
FILE REFERENCE: 01/22080
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Barnea, Eilon
APPLICANT: Beer, Ilan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT
ORGANISM: Artificial Sequence
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Ziv, Tamar
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   38.6%;
100.0%;
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71.4%;
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Pred. No. 3.7e
1; Mismatches
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Score 22;
Pred. No.
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Pred. No. 3
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3.7e+02;
 DB 9; L
               Length 9;
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GENERAL INFORMATION:

GENERAL INFORMATION: MART-1 COMPOUNDS FOR THERAPY AND TITLE OF INVENTION: MART-1 COMPOUNDS FOR THERAPY AND TITLE OF INVENTION: DIAGNOSIS AND METHODS FOR USING SAN FILE REFERENCE: GZ 2103.20

CURRENT APPLICATION NUMBER: US/10/066,474

CURRENT FILING DATE: 2002-01-31

PRIOR APPLICATION NUMBER: US 09/922,405

PRIOR FILING DATE: 2001-03-01

PRIOR FILING DATE: 2001-03-04

PRIOR APPLICATION NUMBER: US 60/23,641

PRIOR APPLICATION NUMBER: US 60/255,502

PRIOR APPLICATION NUMBER: US 60/264,432

PRIOR APPLICATION NUMBER: US 60/264,432

PRIOR APPLICATION NUMBER: US 60/264,432

PRIOR APPLICATION NUMBER: US 60/279,005

PRIOR APPLICATION NUMBER: US 60/279,005

PRIOR APPLICATION NUMBER: US 60/279,005
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US-08-424-550B-278
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US-10-066-474-23
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Best Local Similarity
Matches 4; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Homo sapien
ZIP: 60064-3500
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                              NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
                                                                                                                                                                    IPPLICANI: LU. NON-A, NON-A TITLE OF INVENTION: REAGENTS AND TITLE OF INVENTION: REAGENTS AND TITLE OF SECULENCES: 716
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                                                                      COUNTRY:
                                                                                                              CITY: ABBOTT PARK
                                                                                                                             STREET:
                                                                                                                                               ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                  78, Application US/08424550B
US20020119447A1
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                                                                                                                                                                                                                              SURESH M. DESAI
THOWAS P. LEARY
ANTHONY SCOTT MUERHOFF
JAMES C. ERKER
SHERI L. BUIJK
ISA K. MUSHAHWAR
                                                                                                                             E: ABBOTT LABORATORIES
100 ABBOTT PARK ROAD
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No. US20030013075A1
                                                                          USA
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GEORGE G. SCHLAUDER
                                                                                                                                                                                                                                                                                                                                                                               JOHN N. SIMONS
TAMI J. PILOT-MATIAS
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; Pred. No.
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:

APPLICATION NUMBER:

US/08/424,550B

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                                                                                                                                                                             RESULT 15
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                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: F
SEQ ID NO 9
LENGTH: 11
                                                                                                        Sequence 9, Application US/09845849
Patent No. US20020029395A1
GENERAL INFORMATION:
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Best Local Similarity
Matches 4; Conser
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Best Local Similarity 66.
Matches 4; Conservative
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Patent No. US20010049831A1
GENERAL INFORMATION:
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                                                       APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/291,809C CURRENT FILING DATE: 1999-04-14 PRIOR APPLICATION NUMBER: PCT/US99/08151 PRIOR FILING DATE: 1999-04-13
APPLICANT: WEIGEL, Detlef
APPLICANT: WEIGEL, Igor
APPLICANT: KARDAILSKY, Igor
TITLE OF INVENTION: FLOWERING LOCUS T (FT) AND GENETICALLY
TITLE OF INVENTION: MODIFIED PLANTS HAVING MODULATED FLOWER DEVELOPMENT
                                                                                      APPLICANT: THE SALK INSTITUTE FOR BIOLOGICAL STUDIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 09/060,726 PRIOR FILING DATE: 1998-04-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: FLOWERING LOCUS T (FT)
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APPLICANT: Salk Institut
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LENGTH: 10 amino acids
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REGISTRATION NUMBER: 33,207
REFERENCE/DOCKET NUMBER: 5527.PC.01
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: 435435
ATTORNEY/AGENT INFORMATION:
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GY: linear ·
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66.7%;
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MODULATED FLOWER DEVELOPMENT
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US-08-195-006-40
                US-09-139-762A-74
US-08-469-141A-15
US-08-469-141A-51
US-08-469-141A-57
PCT-US95-13794-51
PCT-US95-13794-51
PCT-US95-13794-42
PCT-US95-13794-42
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Sequence 4, Appli	Sequence 4, Appli	Sequence 4, Appli	Sequence 35, Appl	Sequence 39, Appl	Sequence 9, Appli	Sequence 39, Appl	Sequence 39, Appl	Sequence 37, Appl	Sequence 278, App	Sequence 517, App	Patent No. 5231168	Sequence 38, Appl	. Sequence 7, Appli	Sequence 36, Appl	Sequence 7, Appli	Sequence 7, Appli	Sequence 48, Appl

ALIGNMENTS

RESULT 1 US-08-448-547-2 STRANDEDNESS: TOPOLOGY: linear MOLECULE TYPE: peptide US-08-448-547-2 Sequence 2, Application Patent No. 5821068 GENERAL INFORMATION: COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PATENTIN Release #1.0, Ver. CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/448,547 FILING DATE: 30-UN-1995 CLASSIFICATION DATA: APPLICATION NUMBER: PCT/JP94/01844 APPLICATION NUMBER: PCT/JP94/01844 PRIOR APPLICATION DATA: APPLICATION NUMBER: PCT/JP94/01844 PRIOR APPLICATION DATA: APPLICATION NUMBER: JP 5-297325 FILING DATE: 01-NOV-1993 PRIOR APPLICATION DATA: APPLICATION NUMBER: JP 5-297325 FILING DATE: 02-NOV-1993 TELEFAX: (202)293-7860 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: FILING DATE: 02-NOV-1993 ATTORNEY/AGENT INFORMATION: NAME: Nakamura, Dean H. REGISTRATION NUMBER: 33,981 REFERENCE/DOCKET NUMBER: Q-38 TELECOMMUNICATION INFORMATION: TELEPHONE: (202)293-7060 APPLICANT: APPLICANT: NUMBER OF SEQUENCES: 3 CORRESPONDENCE ADDRESS: APPLICANT: Ito, Yumiko TITLE OF INVENTION: ANTI-HUMAN SOLUBLE FIBRIN ANTIBODY, TITLE OF INVENTION: HYBRIDOWA, AND IMMUNOASSAYING METHOD APPLICANT: STATE: D.C. COUNTRY: USA STREET: 2100 Peni CITY: Washington LENGTH: ADDRESSEE: 2, Application US/08448547 5, 5821068 amino acid Soe, Gilbu Kohno, Isao Inuzuka, Kimiko Ito, Yumiko 3: SUGHRUE, MION, ZINN, MACPEAK & SEAS 2100 Pennsylvania Avenue, N.W. 10 amino (202) 293-7860 acids Release #1.0, Version 2 Q-38931 #1.30

Best Local

Similarity

93.0%; Score 53; 100.0%; Pred. No.

DB 2; L

Length 10;

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Query Match
Best Local Similarity
Atches 8; Conserve
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 9
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NAME: GIDIII, James A.
REGISTRATION NUMBER: 25772
REFERENCE/DOCKET NUMBER: MS!
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OPERATING SYSTEM: DOS
SOFTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.44Mb Storage
                                                                                                                                                                                                                                                                   DESCRIPTION: peptide PUBLICATION INFORMATION:
                                                                                                                                                                                                                                                                                                 MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: Bayer Corporation
STREET: 800 Dwight Way
STREET: P. O. Box 1986
CITY: Berkeley
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                                                                                                                                                                                                                                                                                                                             STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
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                                                                                                                                                                                                            AUTHORS:
                                                                                                                                                                                                                                                                                                                    TOPOLOGY:
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                                                                                      28-JUN-1990
                                                                                               Hader, P.
Walder, P.
Straub, P. W.
Straub, P. W.
Isolation of Human Fibrinogen and its
Derivatives by Affinity Chromatography on Gly-Pro-
Arg-Pro-Lys-Fractogel
Thromb. Haemost.
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Haeberli, A.
                                                                                                                                                                                                                                                                                                                                                                                     (510)705-7904
(570) TD NO: 5:
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          Conservative
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          Score 45; DB 1; Le
Pred. No. 1.9e+05;
1; Mismatches 0;
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SOFTWARE: Patentin Ve
SEQ ID NO 10
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GENERAL INFORMATION:
APPLICANT: Nycomed Imaging AS
TITLE OF INVENTION: Improvement
TITLE OF INVENTION: Agents
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/08/958,993A
CURRENT FILING DATE: 1997-10-28
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CURRENT APPLICATION NUMBER: US/08/960,054A
CURRENT FILING DATE: 1997-10-29
NUMBER OF SEQ ID NOS: 31
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TITLE OF INVENTION:
TITLE OF INVENTION:
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                                                                                                                               TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                              OTHER INFORMATION:
                                                               NAME/KEY: MOD_RES
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OTHER INFORMATION:
NAME/KEY: MOD_RES
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ORGANISM: Artificial Sequence
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INFORMATION: AMIDATION
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                                              Biotinylated-Gly
                                                                                              Description of Artificial Sequence:Biotinylated fibrin-anti-polymerant peptide
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: 31
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: REF/Klaveness/206
CURRENT APPLICATION NUMBER: US/08/959,206A
CURRENT FILING DATE: 1997-10-24
NUMBER OF SEQ ID NOS: 27
                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                      itent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LOCATION: (1)
OTHER INFORMATION: Biotin-Gly
NAME/KEY: MOD RES
LOCATION: (10)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: AMIDATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: Description of Artificial Sequence:Biotinylated OTHER INFORMATION: fibrin-antipolymerant peptide NAME/KEY: MOD_RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Artificial Sequence
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/762,695
FILING DATE: 12-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
                                                                                                                                                                                                                                                           APPLICANT: VON DER ELTZ, HERBERT
TITLE OF INVENTION: SYNTHETIC STANDARD FOR IMMUNOASSAYS
NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                              NERAL INFORMATION SEIDEL, CHRIS
                                                                                                                                                                                                          ADDRESSEE: NIKAIDO, MARMELSTEIN, MURRAY, AND ORAM STREET: METROPOLITAN SQUARE, 655 15TH ST, N.W., SUITE STREET: 330-G ST.LOBBY
                                                                                                                                                                        CITY: WASHINGTON STATE: D.C.
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                                                                                                                                                         COUNTRY: USA
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8; Conserv
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                                                                                                                                         20005-5701
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5846738
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8; Conserv
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80.0%;
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Pred. No. 0.054;
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Pred. No. 0.054;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 10;
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; TOPOLOGY: 1i
; MOLECULE TYPE:
US-08-762-695-11
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: DI
FILING DATE: 20-MAY-11
PRIOR APPLICATION DATA:
APPLICATION NUMBER: DI
FILING DATE: 15-JUN-12
TELEFAX: (415) 326-2422 INFORMATION FOR SEQ ID NO:
                      NAME: Dow, Karen B.
REGISTRATION NUMBER: 29,684
REFERENCE/DOCKET NUMBER: 12
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 326-2400
                                                                                                            CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/089,455
FILLING DATE: 09-JUL-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                 ZIP: 94301
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Lanza, Francois
APPLICANT: Phillips, David R.
APPLICANT: Cazenave, Jean-Pierre
TITLE OF INVENTION: Platelet Glycoprotein V Gene and Uses
NUMBER OF SEQUENCES: 43
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: NOLAN, SHARON L.
REGISTRATION NUMBER: 36,;
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                     CURRENT APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
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TYPE: amino acid
STRANDEDNESS: single
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STATE: California
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                                                                                                                                                                                                                                                                                                                                                                                                       379 Lytton Avenue
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Pred. No.
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RESULT 8
US-08-195-006-40
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Best Local Similarity
Matches 7; Conserv:
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GENERAL INFORMATION:
APPLICANT: Lanza, F
APPLICANT: Phillips
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                                                                                                              TELEFAX: (415) 326-242
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                    TYPE: amino acid
TOPOLOGY: unknown
MOLECULE TYPE: pepi
                                                                                                                                                          TELECOMMUNICATION INFORMATION: TELEPHONE: (415) 326-2400
                                                                                                                                                                                                                                    APPLICATION NUMBER: US 08/089,455
FILING DATE: 09-JUL-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                     HYPOTHETI ÇAL:
                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
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ADDRESSEE: Townsend and Townsend Khourie and Crew
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OTHER INFORMATION:
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            NAME/KEY:
                                                                                                                                                                                   NAME: Dow, Karen B.
REGISTRATION NUMBER: 29,684
REGISTRATION NUMBER: 12418-28
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                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/0:
FILING DATE: 10-FEB-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Palo Alto
STATE: California
                                                                                                                                                                                                                                                                                                  CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
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                                                                                                   ENGTH:
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o. 6083688
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Peptide
                                                                                                                                            (415) 326-2422
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Phillips, David R.
Cazenave, Jean-Pierre
VENTION: Platelet Glycoprotein V Gene and Uses
                                                                   unknown
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                                       :: peptide
NO
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identical to GPV."
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                                                                                                                               40:
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Pred. No.
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PCT-US94-07644A-40
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   Matches
                              Query Match
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Matches
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                                                                                                                                                                                                                                                                                                                TELEFAX: (415) 326-242
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
APPLICATION NUMBER: PC
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy (
COMPUTER: IBM PC com)
                                                                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER: 29,684
REFERENCE/DOCKET NUMBER: 01
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
NAME: Dow, Karen B.
                                                                                                                                                                                                                                                                OLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: COR Therapeutics, Inc.
TITLE OF INVENTION: Platelet Glycoprotein V Gene and Uses
NUMBER OF SEQUENCES: 43
                                                                                                                                                                                                                                                YPOTHETICAL:
Local Similarity nes 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                           OTHER INFORMATION: OTHER INFORMATION:
                                                                                                          NAME/KEY:
LOCATION:
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OTHER INFORMATION:
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OTHER INFORMATION:
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OTHER INFORMATION:
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                                                                                                                                                     OTHER INFORMATION:
                                                                                                                                                                                                    NAME/KEY: Peptide
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les 7; Conserv
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OTHER INFORMATION:
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                                                                                                                                                                                                                                                                                           amino acid
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 Conservative
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            63.2%; Score 36; DB 5; 100.0%; Pred. No. 0.55;
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                                                                                                                                                  /note= "Amino acid sequence of the human fibrinogen (Fg) A-alpha 2 chain thrombin cleavage site."
                                                                      /note= "Amino acid residues identical to GPV."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note= "Amino acid residues
identical to GPV."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note= "Amino acid sequence of the human fibrinogen (Fg) A-alpha 2 chain thrombin cleavage site."
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Pred. No.
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                           Length 11;
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Best Local Similarity
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                                                                                                                                                                                                             -08-441-943-34
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CLASSIFICATION DATA:
PRIOR APPLICATION NUMBER: US 08/7
APPLICATION NUMBER: US 08/7
APPLICATION NUMBER: US 08/7
APPLICATION NUMBER: US 08/7
                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                           APPLICANT:
                                                                                                                                                                            cent No.
                                                             APPLICANT: BURKE, RAE LYN
APPLICANT: RASNUSSEN, MIRELLA EZBAN
APPLICANT: MIKKELSON, JAN MOLLER
TITLE OF INVENTION: PROTEIN COMPLEXES HAVING FACTOR VIII:C
TITLE OF INVENTION: ACTIVITY AND PRODUCTION THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER: CV
TELECOMMUNICATION INFORMATION:
TELEPHONE: 908-281-2372
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APPLICATION NUMBER: US
                               CORRESPONDENCE ADDRESS
                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 02-MAY-1994
ATTORNEY/AGENT INFORMATION:
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                   DDRESSEE:
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E: REED & ROBINS
285 HAMILTON AVENUE, SUITE 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   8 amino acids
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                                                                                                                                             CHAPMAN, BARBARA
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100.0%; Pred. No.
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PRIOR APPLICATION DATA:
RAPPLICATION NUMBER: US 051,916
FILING DATE: 19-MAY-1987
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 822,989
FILING DATE: 27-JAN-1986
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                                         ZIP: 07065-0907

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/441,943
FILING DATE: 16-MAY-1995
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MEDIUM TYPE: Flappy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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CORRESPONDENCE ADDRESS:
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TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                   STREET: 126 I
CITY: RAHWAY
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                                                                                                                                                                                                                    ADDRESSEE: DR. CHRISTINE E. CARTY
STREET: 126 E. LINCOLN AVENUE., P.O. BOX 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REGISTRATION NUMBER:
CLASSIFICATION:
              APPLICATION NUMBER: US/08/469,141A FILING DATE: 06-JUN-1995
                                                                                                                                                                   COUNTRY:
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UNITED STATES OF AMERICA
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                                                                                                                                                                                                                                                                                                                                                                            MUMFORD, RICHARD A. DAVIES, D.T. PHILIP
                                                                                                                                                                                                                                                                                                                          BOGER, JOSHUA S. HUMES, JOHN L.
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POLYMORPHONUCLEAR LEUKOCYTE ELASTASE ACTIVITY
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Pred. No. 1.9e+05;
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                                                                                                           TELECOMMUNICATION INFORMATION:
TELEPHONE: (908) 594-6734
TELEFAX: (908) 594-4720
INFORMATION FOR SEQ ID NO: 33:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
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                      TOPULCE. POMOLECULE TYPE: POMOLECULE TYPE: POMOLECULE TYPE: NO
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FRAGMENT TYPE: internal
                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: Carty, Christine E.
REGISTRATION NUMBER: 36,0
                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: PCT/US95/13794 FILING DATE: 03-NOV-1995
                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
COPERATING SYSTEM: PC-DOS/MS-DOS
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FRAGMENT TYPE:
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HYPOTHETICAL: N
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TRANDEDNESS: single
OPOLOGY: linear
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126 E. Lincoln Avenue; P.O.
                                                                                                                                                                                                                                                                                                                                                                                                     USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Dahlgren, Mary E. Boger, Joshua S. Humes, John L.
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Pred. No.
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                                                Matches
                                                       Query Match
Best Local Similarity
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Best Local
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                        TOPOLOGY: unknown
MOLECULE TYPE: peptide
                                                                                                                                                                                                                        FILING DATE: 08-NOV-
                                                                                                                                                                                                                                            APPLICATION NUMBER: GB 9422534.9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                tent No.
                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
FILING DATE: 02-JUN-1997
                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 9514698.1
                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: Word Perfect CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                    STRANDEDNESS
                                                                                                                                                                                                 APPLICATION NUMBER: GB 9
FILING DATE: 20-AUG-1994
                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: FILING DATE: 17-AUC
                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/09/139,762A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: 1100 New CITY: Washington
                                                                                                                                                              ENGTH:
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                                                                                                                                                             10 amino acids
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1100 New York Avenue, N.W.
                                               Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Klug, Aaron
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         IBM PC compatible
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                                                          47.4%;
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                                           Score 27; DB 3; Pred. No. 28; 3; Mismatches
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Pred. No.
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RESULT 15 US-08-469-141A-15 ; Sequence 15, Application US/08469141A

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Search completed: June 27, 2003, 17:34:50 Job time: 27 secs
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                                                                                                                                                                               Query Match 45.6
Best Local Similarity 100
Matches 5; Conservative
                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFOMMATION:
NAME: CARTY, CHRISTINE E.
REGISTRATION NUMBER: 36,099
REFERENCE/DOCKET NUMBER: 174611B
TELECOMMUNICATION INFORMATION:
TELEPHONE: (908)-594-6734
TELEFAX: (908)-594-4720
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
TYPE: molecular TYPE: peptide
HYPOTHETICAL: NO
ANTI-SENSE: NO
FRAMMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: USA

ZIP: 07065-0907

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION NUMBER: US/08/469,141A
FILING DATE: 06-UN-1995
CLASSIFICATION: 435
CCLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Patent No. 6124107
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: DR. CHRISTINE E. CARTY
STREET: 126 E. LINCOLN AVENUE., P.O. BOX 2000
CITY: RAHWAY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: MUMFORD, RICHARD A.
APPLICANT: DAVIES, D.T. PHILIP
APPLICANT: DAHLGREN, MARY E.
APPLICANT: BOGER, JOSHUA S.
APPLICANT: HUMES, JOHN L.
TITLE OF INVENTION: ASSAY FOR MARKER OF HUMAN
TITLE OF INVENTION: POLYMORPHONUCLEAR LEUKOCYTE ELASTASE ACTIVITY
NUMBER OF SEQUENCES: 71
                                                                                                                                1 GPRVV 5
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100.0%; Pred. No. 1.9e+05;
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Result
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Maximum Match 100%
Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence:
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Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Database :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Searched:
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        Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score
        PIR_73:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*
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Gapop 10.0 , Gapext 0.
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Copyright (c) 1993 - 2003 Compugen Ltd.
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pyruvate dehydroge collagen alpha 1(I Ig heavy chain CRD Ig gamma-2b chain collagen alpha 1(I collagen alpha 1(I collagen alpha 1(I neuropeptide calla glutathione S-tran ribosomal protein phage antigenic de histidine-rich protein collagen alpha I rityus gamma-MSH-like progamma-MSH-like pro
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DNA-binding protei
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3 RPLDKKR 9

RESULT 2 JQ0395 JQ0395 JQ0395 JQ0395 N;Alternate names: hypothetical 1.4K protein C;Species: Azorhizobium caulinodans A;Note: host Sesbania rostrata C;Date: 07-Sep-1990 #sequence_revision 07-Sep-1990 #text_change 03-Feb C;Accession: JQ0395 R;Goethals, K.; Gao, M.; Tomekpe, K.; Van Montagu, M.; Holsters, M. Mol. Gen. Genet. 219, 289-298, 1989 A;Title: Common nodABC genes in Nod locus 1 of Azorhizobium caulinodan A;Reference number: JQ0393; MUID:90136519; PMID:2615763 A;Kolecule type: DNA A;Residues: 1-11 <goe> A;Cross-references: GB:L18897 A;Experimental source: strain ORS571 Query Match Best Local Similarity 71.4%; Pred. No. 1e+02; Matches 5; Conservative 1; Mismatches 1; Indels 0; G</goe>	RESULT 1 S39437 D-amino-acid oxidase (EC 1.4.3.3) - Trigonopsis variabilis (fragment) C;Species: Trigonopsis variabilis C;Date: 19-Mar-1997 #sequence_revision 05-Dec-1997 #text_change 07-May C;Accession: S39437 R;Schraeder, T.; Andreesen, J.R. Eur. J. Biochem. 218, 735-744, 1993 A;Title: Evidence for the functional importance of Cys298 in D-amino a A;Reference number: S39437; MUID:94094869; PMID:7903639 A;Accession: S39437 A;Molecule type: protein A;Residues: 1-9 <sch> A;Experimental source: CBS 4095 C;Function: A;Note: reoxidation of the enzyme by molecular oxygen is accompanied by C;Keywords: FAD; oxidoreductase; catalyzes the oxidation of D-amino acid A;Note: reoxidation of the enzyme by molecular oxygen is accompanied by C;Keywords: FAD; oxidoreductase CyKeywords: FAD; oxidoreductase Ouery Match Best Local Similarity 100.0%; Pred. No. 2.8e+05; Matches 4; Conservative 0; Mismatches 0; Indels 0; G Oy 1 GHRP 4 Db 3 GHRP 6</sch>	30 18 35.3 12 2 PN0046 ATP synthas 31 18 35.3 13 2 E39778 lactose pho 32 18 35.3 13 2 I84603 deoxynuclee 35.3 15 2 PS0212 2 PK proteir 34 18 35.3 16 2 G45681 ribosomal ribosoma
0 2 2	(fragment) 1 D-amino acid oxidase from 2 amino acids to their corres, companied by the release of open companied by the release of	ATP synthase D.cha lactose phosphotra deoxynucleotidyltr 29K protein 4228 - orf 61.1 - phage T ribosomal protein cystic fibrosis tr gene CFTR protein T cell receptor al T cell receptor al hydrogenase (EC 1. cytochrome c (EDH)) 36K microfibril-as hemoglobin alpha c transcription fact rhodopsin homolog

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C;Species: Gallus gallus (chicken)
C;Date: 24-Jan-1992 #sequence_revision
C;Accession: A37968; A36345
R;de la Rosa, E.J.; Kayyem, J.F.; Roman, J.M.; Stierhof, Y.D.; J. Cell Biol. 111, 3087-3096, 1990
A;Title: Topologically restricted appearance in the developing A;Reference number: A36345; MUID:91100421; PMID:2269667
                                                                           A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-19 <DEL>
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                                                                                                                                                                         R;de la Rosa, E.J.; Kayyem,
J. Cell Biol. 112, 1049, 19
A;Reference number: A37968;
                                                                                                                                                                                                                                                                                          neural surface protein Bravo - chicken (fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    submitted to the Protein Sequence A; Reference number: A59069
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                                                                                                                                        A; Accession: A37968
                                                                                                                                                            A;Contents: erratum
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Status: preliminary
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Accession: A59069
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Title: A conserved TATA-less proximal promoter drives; Title: A conserved TATA-less proximal promoter drives; Reference number: I52614; MUID:95329719; PMID:7605992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Accession: I52614
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cross-references: GB:S78532; NID:g999307; PIDN:AAD14289.1; PID:g4261989
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens (man)
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80.0%;
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                                                       J.F.; Roman, J.M.; Stierhof, Y.D.; Dreyer, W.J.; Schwarz, U
                                                                                                                                                                           MUID:91154309;
                                                                                                                                                                                                             J.F.; Roman, J.M.; Stierhof,
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Pred. No
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Pred. No. 4.3e+02;
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                                                                                                                                                                           PMID:1999455
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                       chick retinotectal
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hypothetical EWSR1/FLI1 mutant fusion protein, type 1 - human (fragment) C;Species: Homo sapiens (man) C;Species: Homo sapiens (man) C;Patte: 21-Feb-1997 #sequence_revision 21-Feb-1997 #text_change 20-Apr-2000 C;Accession: I51887 C;Accession: I51887 R;Downing, J.R.; Head, D.R.; Parham, D.M.; Douglass, E.C.; Hulshof, M.G.; Li Am. J. Pathol. 143, 1294-1300, 1993
                                                                                                               RESULT
I51887
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   gene p10 protein - Choristoneura fumiferana nuclear polyhedrosis virus (fragment)
C;Species: Choristoneura fumiferana nuclear polyhedrosis virus, CfMNPV
C;Date: 19-Mar-1997 #sequence_revision 24-Mar-1999 #text_change 24-Mar-1999
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A;Molecule type: protein
A;Residues: 1-7,9-19 <DE
                                                                                                                                                                                                                                                                                                                                       C; Keywords: mitochondrion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Experimental source:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        R;Hill, J.E.; Kuzio, J.; Wilson, J.A.; MacKinnon, E.A.; Faulkner, P. Biochim. Biophys. Acta 1172, 187-189, 1993
A;Title: Nucleotide sequence of the p74 gene of a baculovirus pathogenic to the spruce A;Reference number: S29849; MUID:93176808; PMID:8439559
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A;Residues: 1-12 <TIT>
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A;Residues: 1-12 <HIL>
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R;Hill, J.E.; Kuzio,
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                                                                                                                                                                                                                                                                                                                                                      ;Genome: mitochondrion ;Genetic code: SGC6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ;Species: mitochondrion Crithidia fasciculata
;Date: 30-Sep-1991 #sequence_revision 30-Sep-1991
;Accession: C34135
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Best Local Similarity
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66.7%;
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Pred. No. 7e+0
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7e+02;
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                     Hulshof, M.G.; Link,
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melanotropin beta -
C;Species: Scyliorhi
C;Date: 13-Jul-1981
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                                                                                                                                                                                                          R;Li, Y.; Lacerda, D.A.; Warman, M.L.; Beier, D.R.; Yoshio; Taylor, B.A.; Seegmiller, R.E.; Olsen, B.R. Cell 80, 423-430, 1995
Cell 80, 423-430, 1995
A;Title: A fibrillar collagen gene, Collial, is essential A;Reference number: A55648; MUID:95163095; PMID:7859283
A;Accession: A55648
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C; Superfamil C; Keywords:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gen. Comp. Endocrinol. 24, 398-404, 1974
A;Title: A beta-MSH in the pituitary gland of the spotted dogfish
A;Reference number: A01470; MUID:75113445; PMID:4452470
A;Accession: A01470
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C;Species: Scyliorhinus canicula (smaller spotted catshark, smaller spotted C;Date: 13-Jul-1981 #sequence_revision 13-Jul-1981 #text_change 23-Aug-1996 C;Accession: A01470  
R;Love, R.M., Pickering, B.T.
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C; Keywords: fusion ----
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A; Residues: 1-17 < DOW>
A; Cross-references: GB: 566911; NID: g440935;
C; Comment: This sequence is the chimeric pro
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                                                                                                                                                                                                                                                                                                                                                                     collagen alpha 1(XI) chain - mouse (fragment)
                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 10
                                                                                                                                                                                                                                                                                                                 ;Species: Mus musculus (house mouse);Date: 23-Mar-1995 #text_change 17-Mar-1999;Accession: A55648
                                                                                                                                        Superfamily: collagen
                                                                                                                                                          Molecule type: DNA
Residues: 1-18 <LIA>
                                                                                                                                                                                          Status: preliminary
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Best Local
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KPLDR 11
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45.5%;
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                                                                    Score 21; DB 2;
Pred. No. 1.1e+03;
2; Mismatches 0
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Pred. No.
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Pred. No. 1e+03;
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C;Accession: A23694
R;Cole, D.G.; Yount, R.G.
J. Biol Chem. 265, 22537-22546, 1990
J. Biol Chem. 265, 22537-22546, 1990
A;Title: Photolabeling of the 6 and 10 S conformations of A;Reference number: A23694; MUID:91093105; PMID:2266144
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Molecule type: protein
A; Residues: 1-9 <BIN>
A; Experimental source: human bile and
C; Superfamily: translation elongation
C; Keywords: GTP binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R)Binette, J.P.; Binette, M.B. submitted to the Protein Sequence Database, October 1996 submitted to the Proteins of kidney and gallbladder stones.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R;East, A.K.; Stacey, J.M.; Collins, M.D. Syst. Appl. Microbiol. 17, 306-312, 1994
A;Title: Cloning and sequencing of a hemagglutinin A;Reference number: S58855
A;Accession: S58855
                                                 A;Residues: 1-13 <COL>
C;Keywords: smooth muscle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       translation elongation factor EF-Tu - unidentified C;Species: unidentified bacterium C;Date: 07-Feb-1997 #sequence_revision 07-Feb-1997
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C; Date: 15-Feb-1996 #sequence_revision
C; Accession: $58855
                                                                   A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-13 <COL>
                                                                                                                   A; Reference number: A; Accession: A23694
                                                                                                                                                                                                                                                        myosin heavy chain, smooth muscle -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Accession: D58503
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A;Residues: 1-18 <EAS>
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Pred. No. 1.1e
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Pred. No. 2.8e+05;
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PL0143
Search completed: June 27, Job time : 40 secs
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A;Residues: 1-15 <KRA>
C;Comment: Carbon-monoxide dehydrogenase consists of three polypeptide chains: large,
C;Keywords: oxidoreductase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       carbon-monoxide dehydrogenase (EC 1.2.99.2) medium chain - Pseudomonas carboxydohydrogen C;Species: Pseudomonas carboxydohydrogena C;Datec: 07-Sep-1990 #sequence_revision 07-Sep-1990 #text_change 28-Apr-1993
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                                                                                                                                                                                                                                                                                                                                                                                         R;Kraut, M.; Hugendieck, I.; Herwig, S.; Meyer, O.
Arch. Microbiol. 152, 335-341, 1989
A;Title: Homology and distribution of CO dehydrogenase structural genes in carboxydotrop
A;Reference number: PL0138; MUID:90055678; PMID:2818128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Molecule type: protein
A;Residues: 1-13 <SZU>
A;Experimental source: liver
C;Complex: The enzyme appears to be a heterodimer of alpha and beta chains.
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A;Reference number: A47415; MUID:93352609; PMID:7688733
A;Accession: B47415
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3; Conservative 0; Mismatches 0;
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    SwissProt_40:*
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Q46228 clover prol
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P43172 ascaris suu
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-!- FUNCTION: PROTEIN S19 FORMS A COMPLEX WITH S13 THAT BINDS STR
TO THE 16S RIBOSOMAL RNA (BY SIMILARITY).
-!- SIMILARITY: BELONGS TO THE S19P FAMILY OF RIBOSOMAL PROTEINS.
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-!- FUNCTION: PROTEIN S19 FORMS A COMPLEX WITH S13 THAT BINDS STRONGLY
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-!- SIMILARITY: BELONGS TO THE S19P FAMILY OF RIBOSOMAL PROTEINS
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Bacteria; Firmicutes; Mollicutes; Acholeplasmatales;
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-!- FUNCTION: PROTEIN S19 FORMS A COMPLEX WITH S13 THAT BINDS STF.
TO THE 16S RIBOSOMAL RNA (BY SIMILARITY).
-!- SIMILARITY: BELONGS TO THE S19P FAMILY OF RIBOSOMAL PROTEINS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Entropean Bioinformatics Institute. There are no restructe by non-profit institutions as long as its content modified and this statement is not removed. Usage by and
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-i- FUNCTION: PROTEIN S19 FORMS A COMPLEX WITH S13 THAT TO THE 16S RIBOSOWAL RNA (BY SIMILARITY).
-i- SIMILARITY: BELONGS TO THE S19P FAMILY OF RIBOSOMAL
                                                                                                  the European Bioinformatics Institute. The use by non-profit institutions as long modified and this statement is not removed. entities requires a license agreement (See
                                                                                                                                                                                                                                               J. Bacteriol. 176:5244-5254(1994).
-!- FUNCTION: PROTEIN S19 FORMS A COMPLEX WITH S13 THAT BINDS STITO THE 16S RIBOSOMAL RNA (BY SIMILARITY).
-!- SIMILARITY: BELONGS TO THE S19P FAMILY OF RIBOSOMAL PROTEINS.
                                                                                                                                                                                                                                                                                                                                       MEDLINE=94350802; PubMed=8071198;
Gundersen D.E., Lee I.M., Rehner S.A.,
"Phylogeny of mycoplasmalike organisms
their classification.";
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between the Swiss Institute of Bioinformatics and the EMBL
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NCBI_TaxID=35776;
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30S ribosomal
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an email to license@isb-sib.ch).
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                     IPR002222;
                                                                                   email to license@isb-sib.ch).
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se I.M., Rehner
RIBOSOMAL_S19;
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(Scyliorhinus Gen. Comp. End.
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21-JUL-1986
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SEQUENCE
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15-JUL-1999 (Rel. 34, Last sequence up
15-JUL-1999 (Rel. 38, Last annotation
Unknown protein from 2D-page of etiola
(Fragment).
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P80623;
01-OCT-1996
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Pernollet J.-C., Zivy M., de Vi
"The maize two dimensional gel
genome analysis program.";
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NON TER 1 1
SEQUENCE 14 AA; 1642 MW; 200
                  Love R.M., Pickering B.T.;

"A beta-MSH in the pituitary gland of the spotted (Scyllorhinus canicula): isolation and structure."

Gen. Comp. Endocrinol. 24:398-404(1974).
                                                                                                                                                                                             Elasmobranchii; Galeomorphii; Galeoidea;
                                                                                                                                                                                                                                      Melanotropin beta (Beta-MSH).
Scyliorhinus canicula (Spotted dogfish)
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Spermatophyta; Magnoliophyta;
Panicoideae; Andropogoneae; Ze
Panicoideae; Andropogoneae; Ze
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                                                                                            SEQUENCE.
MEDLINE=75113445;
                                                                                                                                                            Scyliorhinidae;
NCBI_TaxID=7830
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62.5%;
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yta; Liliopsida; Poales; Poaceae; PACC
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Pred. No. 1.7e+02;
1; Mismatches 0
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Pred. No.
                                                                                                                                                                                               id dogfish) (Spotted catshark).
Craniata; Vertebrata; Chondri
Galeoidea; Carcharhiniformes;
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DE Cerbo
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OC Bacte
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Best Local'S
Matches 5
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01-FEB-1991 (Rel. 17, Last sequence update)
01-TEB-1991 (Rel. 17, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Carbon monoxide dehydrogenase medium chain (EC dehydrogenase subunit M) (CO-DH M) (Fragment).
                                                                                                                                                                                [1]
SEQUENCE.
MEDLINE=90055678; PubMed=2818128;
MEDLINE=9005678; PubMed=2818128;
MEDLINE=9005678; PubMed=281818128;
MEDLINE=9005678; PubMed=281818128;
MEDLINE=9005678; PubMed=281818128;
MEDLINE=9005678; PubMed=28181828;
MEDLINE=9005678; PubMed=28181828;
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P19917;
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P43172;
01-NOV-1995
                                                                                                                                                       carboxydotrophic bacteria.", Arch. Microbiol. 152:335-341
                                                   -!- CATALYTIC ACTIVITY:
                                                                                                                                                                                                                                                                                                                                                                                  Bacteria; Proteobacteria;
Bradyrhizobium group.
                                                                                                                                                                                                                                                                                                                                                                                                             Pseudomonás carboxydohydrogena.
Bacteria; Proteobacteria; alpha
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MOD_RES
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Ascaris suum (Pig roundworm) (Ascaris lumbricoides).
Eukaryota; Metazoa; Nematoda; Chromadorea; Ascaridida; Ascaridoidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=95380362; PubMed=7651904; Cowden C., Stretton A.O.W.;
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01-NOV-1995 (Rel. 32, Last sequence update)
01-FEB-1996 (Rel. 33, Last annotation update)
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NCBI_TaxID=6253;
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1012 MW;
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524F073774176877 CRC64;
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P41839;
01-NOV-1995
01-NOV-1995
16-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                   TISSUE=Hemolymph;

MEDLINE=97362903; PubMed=9219367;

Rees J.A., Moniatte M., Bulet P.;

Rees J.A., Moniatte Septides isolated from "Novel antibacterial peptides isolated from Bombus pascuorum (Hymenoptera, Apoldea).";

Insect Biochem. Mol. Biol. 27:413-422(1997)

Insect Biochem. Mol. Biol. 27:413-422(1997)
SEQUENCE.
SEQUENCE.
TISSUB=Thoracic ganglion, Brain,
MEDLINE=93211980; PubMed=8460157;
Nuve H., Johnsen A.H., Scott A.G
                                                                                                       01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Callatostatin 1 (Leu-callatostatin 1) [Contains: Call
callatostatin 2); Callatostatin 3 (Leu-callatostatin
Calliphora vomitoria (Blue blowfly)
                                                             Muscomorpha; Oest
NCBI_TaxID=27454;
                                                                        Eukaryota; Metazoa; Arthropoda; Mandibulata; Pancrustacea; Insecta; Pterygota; Neoptera; Endopterygota; Diptera; Brac
Muscomorpha; Oestroidea; Calliphoridae; Calliphora.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    P81464;
15-DEC-1998
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                                                                                                                                                                                               CALVO
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APID_I
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Oxidoreductase; Flavoprotein;
NON_TER 15 15
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Insecta; Pterygota; Neoptera; Endopterygota;
Aculeata; Apoidea; Apidae; Bombus.
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3; Conserv
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37, Last sequence up
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100.0%; Pr
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Pred. No. 5e+0
2; Mismatches
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          A.G.,
                                                                                                                                                                                                                                                                                                                         Hemolymph.
CD1D0D02C8BC23D1
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         Yagi K.J.,
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PEPTIDE '
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15-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "Distribution and functional significance of Leu-callatostatins in the blowfly Calliphora vomitoria.";
Cell Tissue Res. 276:367-379(1994).
-i- FUNCTION: MAY ACT AS A NEUROTRANSMITTER OR NEUROMODULATOR AND F A ROLE IN THE INTEGRATION OF INFORMATION WITHIN THE BRAIN. MAY INVOLVED IN THE CONTROL OF VISCERAL MUSCLES DUE TO ITS ABILITY BEHAVE AS POTENT INHIBITORS OF PERISTALIC MOVEMENTS. MAY ALSO FULFILL A NEUROHORMONAL ROLE ON MUSCLES OF THE GUT AND HEART.
-i- TISSUE SPECIFICITY: BRAIN, SUBDESOPHAGEAL GANGLION, RETROCEREBRE CONTROL OF THE GUT AND HEART.
                                                                                                                                                                                                                             MEDLINE=96190713; PubMed=8611151;
Becerril B., Corona M., Coronas F.I., Zamudio F.,
Calderon-Aranda E.S., Fletcher P.L. Jr., Martin B.M.,
"Toxic peptides and genes encoding toxin gamma of the
scorpions Tityus bahiensis and Tityus stigmurus.";
Biochem. J. 313:753-760(1996).
-!- FUNCTION: NOT TOXIC IN MICE.
                                                                                                           Ptam;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Toxin TbTX-VI (Fragment).
Tityus bahiensis (Brazilian scorpion)
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Buthoidea; Buthidae; Tityus.
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15-DEC-1998 (Rel. 37, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PIR;
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"Callatostatins:
                                                                                       SEQUENCE
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R; A47393; A47393.
R; B47393; B47393.
R; C47393; C47393.
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SIMILARITY: BELONGS TO
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Pred. No. 7.
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30-MAY-2000
                                                                                                         01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Melanotropin gamma (Gamma-melanocyte stimulating hormone)
Theromyzon tessulatum (Leech)
MEDLINE=94298944; PubMed=8026574; Salzet M., Wattez C., Bulet P., Malecha J.; "Isolation and structural characterization of a novel peptide to gamma-melanocyte stimulating hormone from the brain of the
                                                                                                                                                                    MLG_THE'
P41989;
                                                                                                                                                                                                                                                                                                                                                                                                                      entities
or send a
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                                                                                        Theromyzon tessulatum (Leech).
Eukaryota; Metazoa; Annelida; Clitellata; Hirudinida; Hirudinea;
Rhynchobdellida; Glossiphoniidae; Theromyzon.
                                                                                                                                                                                                                                                                                                                                                                                                                                           use by non-profit institutions as long modified and this statement is not removed.
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Spiroplasmataceae; Spi
                                              TISSUE=Brain;
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European Bioinformatics Institute. There a
by non-profit institutions as long as i
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                                                                            [_TaxID=13286;
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TE; PS00323; RIBOSOMAL_S19; PARTIAL
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l; Mismatches
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re greater than or equal to the score of the result being printed,
is derived by analysis of the total score distribution.

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51
1 GHRPLDKKR 9
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sp_bacteria:*
sp_fungi:*
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sp_virus:*
sp_vertebrate:*
sp_unclassified:*
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sp_phage:*
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sp_bacteriap:*
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     GenCore version (c) 1993 - 2003
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Compugen Ltd
                  Q9qxp9 mus musculu
Q94106 brassica na
Q9um92 homo sapien
Q8ztx6 pyrobaculum
Q63058 lathraea cl
Q9um87 homo sapien
Q8rjf1 pseudomonas
P83331 streptococc
Q8t8b6 ciona intes
Q9psw5 gallus gall
Q98y97 human immun
Q45866 clostridium
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Q56750 western x p
Q9t2q6 solanum tub
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ALIGNMENTS

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RESULT
Q56750
ID Q5
AC Q5
AC Q5
DT 01
DT 01
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Best Local Similarity
Matches 5; Conserv
   Q56750;
Q56750;
01-NOV-1996
01-NOV-1996
01-DEC-2001
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Q46291;
01-NOV-1996
01-NOV-1996
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01-NOV-1996 (TrEMBLrel. 19, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Ribosomal protein S19 (Fragment)
Canadian peach X phytoplasma
Bacteria; Firmicutes; Bacillus/Clostridium group; Mollicutes;
Acholeplasmatales; Acholeplasmataceae; Phytoplasma.
NCBI_TaxID=35778;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               J. Bacteriol. 176:5244-5254(1994).
EMBL; L27016; AAA83950.1; -.
NON_TER 1 1
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Gundersen D.E., Lee I.M., Rehner S.A.,
"Phylogeny of mycoplasmalike organisms
their classification.";
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   (TrEMBLrel. 01, (TrEMBLrel. 01, (TrEMBLrel. 19,
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62.5%;
Created)
Last sequence update)
Last annotation update)
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Pred. No. 2e+02;
0; Mismatches
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Q83965

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RESULT 4
Q90XP9
ID 090X
AC Q90X
AC Q90X
DT 01-M
DT 01-M
DT 01-M
DE Hepa
GN HHF3
OS Mus
OC Euka
OC Mus
OC Mamm
OX NCBI
RN [1]
RN [1]
RP SEQU
RC STRA
RA Maeyy
RT "Mus
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Best Local S
Matches 4
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Best Local
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01-MAY-2000
01-MAY-2000
                                                  Eukaryota; Metazoa;
Mammalia; Eutheria;
NCBI_TaxID=10090;
  Maeyama K., Yo
                     SEQUENCE FROM N.A.
STRAIN=C57BL/6J;
                                                                                    Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                        Q9T2Q6;
01-MAY-2000
01-MAY-2000
01-MAR-2001
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                                                                                                                                                                                                                                                                                                          MEDLINE=94198758; PubMed=7764624; Braun H.P., Kruft V., Schmitz U.K.; Planta 193:99-106(1994).
                                                                                                          Hepatocyte nuclear
                                                                                                                                                                                                                                                                                                     SEQUENCE
                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Viridiplantae; Streptophyta; Embryophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eu
Asteridae; euasterids I; Solanales; Solanaceae; Solan
NCBI_TaxID=4113;
                                                                                                                                                                                                                                                                                                                                                                                                         Mitochondrion.
                                                                                                                                                                                                                                                                                                                                                                                                                  Solanum tuberosum
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE
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Western X phytoplasma.
Bacteria; Firmicutes; Bacillus/Clostridium group; Nacholeplasmatales; Acholeplasmataceae; Phytoplasma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=94350802; PubMed=8071198;
Gundersen D.E., Lee I.M., Rehner S.
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) (TrEMBLrel. 13,
) (TrEMBLrel. 13,
  Yoshihashi H.,
us (C57BL/6J) H
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(TrEMBLrel. 13, Last sequence update)
(TrEMBLrel. 16, Last annotation update)
reductase 11 kDa subunit (EC 1.10.2.2)
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                                                               Chordata;
Rodentia;
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57.1%;
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 ., Kosaki R.,
HNF3-beta ge
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Pred. No. 2.3e
2; Mismatches
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Pred. No. 2e+0;
0; Mismatches
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                                                            Craniata; Vertebrata; Euteleostomi;
Sciurognathi; Muridae; Murinae; Mus
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 R., Suzuki
gene, prom
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Kosaki K.;
r region.";
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eudicots;
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RESULT 6
Q9UM92
ID Q9UM
AC MAMM
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AC Q9UM
AC Q9
AC Q
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Q94L06
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Best Local
   EMBL; S78
NON TER
SEQUENCE
                                                      from the Blood 86:
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Q9UM92;
01-MAY-2000
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NON_TER
NON_TER
SEQUENCE
                                                                                                        Blasi F.;
                                                                                                                          Soravia
                                                                                                                                     SEQUENCE FROM N.A.
MEDLINE=95329719;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Fourmann M., Froger N., Bruner C., Tools designi "Amplified consensus gene markers: Tools designi of Arabidopsis-known-function genes in Brassica. Submitted (APR-2000) to the EMBL/GenBank/DDBJ designi apprendiction of ARK49052.1; -
                                                                                                                                                                                                               Eukaryota;
Mammalia;
                                                                                                                                                                                                                                                                                                  01-MAY-2000
                                                                                      "A conserved TATA-less
                                                                                                                                                                                       NCBI_TaxID=9606;
                                                                                                                                                                                                                                            Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core everosids II; Brassicales; Brassicaceae; Brassica.
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Eukaryota; Viridiplantae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=3708;
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                                                  the urokinase-type plasminogen 86:624~635(1995).
                                    S78532;
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   18 AA;
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(TrEMBLrel. 19, Last annotation
plate synthase (Fragment)
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                                  AAD14289.1; -.
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                                                                                                                     PubMed=7605992; A., De Luca P.,
   1964 MW;
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Primates;
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01-MAR-2002 (Tremburg: -
01-MAR-2002 (Tremburg: 2)
01-MAR-2002 (Tremburg: 2)
01-MAR-2002 (Tremburg: 2)
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01-AUG-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         063058
                                                                                                                                                                                                                     Chloroplast:

Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;

Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;

Asteridae; euasterids I; Lamiales; Orobanchaceae; Lathraea.
                   Chloroplast.
                                                                                             EMBL;
                                                                                                          Lusson N., Delavault P., Thalouarn P.;
"The rbcL gene from Lathraea (holparasitic) i
plastid-encoded RNA polymerase.";
Submitted (OCT-1997) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                           063058;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "Genome sequence of the hyperthermophilic crenarchaeon aerophilum.";
Proc. Natl. Acad. Sci. U.S.A. 99:984-989(2002).
EMBL; AE009908; AAL64633.1; -.
                                                                                                                                                                                                                                                                                                                                                RPS11
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Archaea; Crenarchaeota;
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 SEQUENCE
                                                   InterPro; IPR001971; Ribosomal_S11
Pfam; PF00411; Ribosomal_S11; 1.
                                                                                                                                                                                        SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                          Lathraea clandestina.
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MEDLINE=96253074; PubMed-
Wada C., Ishigaki M., To,
Yamazaki Y., Ohtani H.;
"[Nucleotide sequences at
growth factor receptor 2
                                                                                                     EMBL; AJ4
NON TER
SEQUENCE
                                                                                                                                                         Submitted (DEC-2001) to the EMBL; AJ422128; CAD19527.1; EMBL; AJ422129; CAD19528.1; EMBL; AJ422130; CAD19529.1;
                                                                                                                                                                                                                                                                                     STRAIN=NC16-2, TC29-5, AND TC97;
TRANSPOSON=TN5041A1, TN5041B, AND TN5041D1;
Kholodii G.Y., Gorlenko Z.M., Mindlin S.Z.,
"Distribution of distinct microvariants of 1
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Bacteria; Proteobacteria;
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Mammalia; Eutheria;
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M., Toyo-oka Y.,
                                                                                                        1206 MW;
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MEDIJNE=21863848; PubMed=11875658;
Takamura K. Egawa T., Ohnishi S., Oka
"Developmental expression of ascidian
genesi. Choline acetyltransferase and
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Eukaryota; Metazoa; Chordata; Urochordata; Ascidiacea; Enterogona;
Phlebobranchia; Cionidae; Ciona.
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01-JUN-2002 (TrEMBLrel. 21, Created)
01-JUN-2002 (TrEMBLrel. 21, Last seque
01-JUN-2002 (TrEMBLrel. 21, Last annot
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01-JUN-2002 (TrEMBLrel 21, Last annotation update)
Penicillin-binding protein (PBP) (Fragment).
Streptococcus thermophilus.
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01-JUN-2002
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KPLEKR 13
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                                                                            Score 22; DB
Pred. No. 1.7e
3; Mismatches
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Pred. No.
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                                                                    DB 5;
1.7e+03;
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neurotransmitter synthesi
acetylcholine transporter
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RESULT 15
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Best Local
Q45866;
Q45866;
01-NOV-1996
01-NOV-1996
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01-JUN-2001
01-JUN-2001
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Q98Y97;
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01-MAY-2000
01-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                            amplification products
Submitted (FEB-2001) to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Viruses; Retroid viruses; NCBI_TaxID=11676;
                                                                                                                                                                                                                                                                                                                                      SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human immunodeficiency virus type 1.
Viruses; Retroid viruses; Retroviridae; Lentivirus.
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Gene 166:261-266(1995)
EMBL; L27342; AAA92641.1; -.
NON TER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Korn K.;
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MEDLINE=96125200; PubMed=8543172;
Krishnan V.A., Schatzle J.D., Hin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Archosauria; Aves; Neognathae; Galliformes; Phasiani
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Gallus gallus (Chicken).
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01-MAY-2000 (TrEMBLrel. 13, Last annotation updat
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Last annotation updat
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Pred. No. 2e+(
0; Mismatches
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1; Mismatches
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Phasianidae; Phasiani
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avian
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(TrEMBLrel.

01,

Created) Last seq

sequence update)

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DT 01-DEC-2001 (TrEMBLrel. 19, Last annotation update)

DE NTNH protein.

GN NTNH.

OS Clostridium botulinum.

OC Clostridiales; Firmicutes; Bacillus/Clostridium group; Clostridia;

OC Clostridiales; Clostridiaceae; Clostridium.

OX NCBI_TaxID=1491;

RN [1]

RN SEQUENCE FROM N.A.

RC STRAIN=178;

RA East A.K., Stacey J.M., Collins M.D.;

RT neurotoxin complex encoded by Clostridium botulinum types A and B.";

RT neurotoxin complex encoded by Clostridium botulinum types A and B.";

RSyst. Appl. Microbiol. 17:306-312(1994).

DR EMBL; X79103; CAA55712.1; -.

SQ SEQUENCE 18 AA; 2029 MW; D4DD49856A10DFD9 CRC64;

Query Match
Best Local Similarity 50.0%; Pred. No. 2.5e+03;

Matches 3; Conservative 2; Mismatches 1; Indels. 0; Gaps 0;

PLDKKR 9

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Db 12 PVDNKK 17

Search completed: June 27, 2003, 17:45:58

Job time: 78 secs
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Maximum DB
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51
1 GHRPLDKKR 9
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1: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1980.DAT:*

2: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1981.DAT:*

3: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1982.DAT:*

4: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1984.DAT:*

5: /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1984.DAT:*

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Gapop 10.0 ,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GenCore version 5.1.6 (c) 1993 - 2003 Compugen Ltd.
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and is derived by analysis of the total score distribution.	score greater than or equal to the score of the result being printed,	Pred. No. is the number of results predicted by chance to have a
distributi	e result b	y chance t
on.	eing printed,	o have a

SUMMARIES

10	9	8	7	o	_ل	.4	ω	2	н	Result No.
41	41	41	41	41	51	51	51	51	51	Score
80.4	80.4	80.4	80.4	80.4	100.0	100.0	100.0	100.0	100.0	Query Match
80	7	7	7	7	16	12	12	10	. 9	Query Match Length DB
9	15	15	11	æ	14	14	13	16	22	DB
AAP82686	AAR65792	AAR63269	AAR05558	AAP71314	AAR44830	AAR44829	AAR28629	AAW11931	ABB76979	ID
Human fibrin beta	Fibrin-specific ep	Beta-peptide used	Peptide antigenic	Sequence of fibrin	Human fibrin beta-	Human fibrin beta-	N-terminal human f	Fibrinogen A-alpha	Residues 45-54 of	Description

MO-MuLV variable	AAU75028	23	y	49.0		U
Membrane arbebaraa	11111111111111111111111111111111111111	0 0	,			1 4
Cortabolished	AAW214/5	3 F	ט ע	•) N	. د
	AAY /6363	, ₁ ,	Ţ	٠		, ,
epti	AAE06674	22	17	51.0	200	, μ
ent o	AAY02812	20	17	٠	26	0
מת	ABB72935	23	15	•		φ
derived	AAB97760	22	15	51.0		ω
derived	AAB97759	22		٠		7
TNF-alpha antagoni	AAB84092	22		٠	26	01
Immune modulating	AAB84079	22	15	51.0	26	٠.
. Interferon-gamma	AAB98150	22	15	٠	26	-
Interferon-gamma	AAB98149	22		•	26	_
9	AAM51531	22			26	
Fragment of human	AAY76276	21	15	•	26	
TNF-antagonist pep	AAB17973	21	15	•	26	_
SV40 large T ant	AAB20990	21	13	•	26	_
Human microtubu	AAY20425	19	12	51.0		_
Human platelet gl:	AAW32704	18	10	٠		7
Thrombin cleavage	AAB37079	21	ហ	•	26	•
	AAW34571	18	S	٠	26	•
Non-crosslinked pr	AAY31030	20	4	٠	26	
Accelerator pep	AAR62426	16	4	٠	26	
F	AAR44726	14	18	52.9	27	
BCL2 im	AAY20168	19	17		27	:
Human peptide #27	AAM97002	22	14	•	27	Ŭ
Multiply branched	AAW65554	19	11	•		_
ğ	AAW65555	19	12	٠	29	~
ü	AAR93400	16	14		30	7
Fibronectin beta	AAB10629	21	11		36	•
Fibrinogen-beta	AAB10666	21	11	70.6	36	
Human fibrin beta	AAR44838	14	12	•	41	_
ř	AAR65793	15	œ	80.4	41	_
etic bet	AAR63270	15	8	80.4	41	
Human fibrin be	AAR44837	14	89	80.4	41	•

ALIGNMENTS

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RESULT 1
ABB76 J
ABB76 J
AB 7 ABB77
XX ABB77
XX ABB7
XX AAL1
XX ABB7
X
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22-JUL-2002
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Anti-arthritic; anti-inflammatory; fibrin; rheumatoid polyarthritis; human; fibrinogen beta chain.

Homo sapiens

FR2795735-A1

05-JAN-2001.

01-JUL-1999; 99FR-0008470

01-JUL-1999; 99FR-0008470

(UYTO-) UNIV TOULOUSE SABATIER PAUL.

Serre G, Sebbag M;

WPI; 2001-114394/13.

New citrulline-containing polypeptide from fibrin, useful for diagnosis and treatment of rheumatoid polyarthritis

Example 1; Page 12; 23pp; French.

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RESULT 2
ADMIT 931
ADMIT 931
XX
ADMIT 931
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ADMIT 931
XX
ADMIT 931
XX
ADMIT 92-1
XX
ADMIT 931
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                                          Que Match
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Matches 9
                Matches
                                                                                                                   Sequence
                                                                                                                                                     soluble fibrin in plasma samples (e.g. by sandw particularly for the diagnosis of pathological disseminated intravascular coagulation.
                                                                                                                                                                                                        A novel monoclonal antibody (MAb) reacts with human soluble fibrin, but not with human fibrinogen. Specifically when the MAb reacts with urea-treated des AABB fibrin the reaction is not inhibited by peptides corresponding to fibrinogen A-alpha-chain residues 17-26 (AAWI1931), B-beta-chain residues 15-24 (AAWI1932) or gamma-chain residues 312-324 (AAWI1933). The MAb is useful in immunoassays for soluble fibrin in plasma samples (e.g. by sandwich immunoassay),
                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 3; Page 15; 32pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Monoclonal antibody reactive with soluble human fibrin - but with fibrinogen, is useful for fibrin immunoassay in plasma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1995-206667/27.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               02-NOV-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-NOV-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11-MAY-1995.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO9512617-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Monoclonal; antibody; human; soluble; fibrin; fibrinogen; urea-treated; des-AABB; A-alpha-chain; immunoassay; diagnosis; disseminated intravascular coagulation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to a citrulline (Cit) containing polypeptide derived from all or part of the alpha- or beta-chains of fibrin by substitution of at least one arginine residue by Cit. The Cit containing polypeptides can be used for in vitro diagnosis of rheumatoid polyarthritis (RP), by detecting disease-specific autoantibodies, and therapeutically for neutralising the RP-associated autoimmune response. The present sequence is a fragment (residues 45-54) of the human fibrinogen beta chain precursor, which was used in an example from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (IATR ) IATRON LAB
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     100.0%; ilarity 100.0%; Conservative 0
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                            Score 51; DB 16;
Pred. No. 0.00058;
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Pred. No.
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Mismatches
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. 0.00058;
thes 0;
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                                                                                                                                                                                       conditions such
                                               Length 10;
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Indels

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Gaps

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XXXXXXX

Human fibrin

beta-chain N-terminal peptide

(1-11)-Cys

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RESULT 3
AAA28629
ID AAR2
XX AAR2
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AC AAR2
XX bisp
CX bisp
XW card
XX peri
XX peri
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XX 19-N
XX 19-N
XX 11-N
XX 11-N
XX 11-N
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AAR44829
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                                                                                                             AAR44829
     20-JUN-1994
                                                          AAR44829;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       was used in the production of bispecific monoclonal antibodies whis are specific for fibrins, but do not bind fibrinogen, and and are specific for anti-thrombolytic substance. The compsn. contg. these absolutes the side effects of prior art Ab targetted thrombolytic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bi:specific antibody useful for treating thrombotic obstructive diseases e.g. cardiac infarction - comprises antithrombus antibody variable region and anti-thrombolytic substance antibody variable region with no heavy chain constant region domains 2 and
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 3; 30pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This sequence represents an N-terminal peptide of human fibrin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1992-383677/47.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Iwasa
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                                                                                                                                                                                                                                                                                                                                               Similarity 100
9; Conservative
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                                                                                                       standard; peptide; 12 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            has enhanced
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                         AA;
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92JP-0055025
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                                                                                                                                                                                                                                                                                                                                                                        Score 51;
Pred. No.
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human fibrin beta-chain N-terminal peptide (1-11)-Cys was synthesised and coupled to BSA for injection into mice. The peptide was used to raise antibodies of human fibrin. Monoclonal antibodies specific for fibrin are used in the production of bispecific monoclonal antibodies which also recognise truncated tPA muteins lacking the finger, EGF and Kringle 1 domains.
                                                                                                                                                                                                                                                                                                        Tissue plasminogen activator; t-PA; mutein; fibrin; antigen; anti-fibrin; monoclonal antibody; hybridoma; thrombolysis antithrombotic agent; bispecific antibody.
                                                  20-JUN-1991;
                                                                                                                                                         19-NOV-1993.
                                                                                                                                                                                                      JP05304992-A.
                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                             Human fibrin beta-chain internal peptide fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAR44830 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hybrid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1993-408334/51.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-JUN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-JUN-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-NOV-1993.
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                                                                                                    17-JUN-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-JUN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      reduced reactivity to fibrinogen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 having
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           monoclonal antibody - used for prepn. of thrombolytic drug increased thrombolytic activity and specificity and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 \vdash
  TAKEDA
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Similarity 100.0%;
9; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GHRPLDKKR
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12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      note= "BSA carrier is"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               peptide;
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Pred. No. 0.00071;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  À
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human fibrin internal peptide fragment was identified as a candidate immunogen to raise antibodies to human fibrin. Monocloi antibodies specific for fibrin are used in the production of bisimonoclonal antibodies which also recognise truncated tPA muteins lacking the finger, EGF and Kringle 1 domains.
                                                                                                                                                                                                                    Screening of fibrin-specific monoclonal antibodies - with immobilised crosslinked fibrin clot and screening
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   03-OCT-2002
19-JUN-1991
                                                                                                                                                    Disclosure; Page 7; 41pp;
                                                                                                                                                                                                                                                                                                                                 Matsueda GR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14-APR-1987;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Misc-difference
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                                                                                                           The MAbs are specific to fibrin without
                                                                                                                                                                                                                                                                                      WPI; 1987-306855/43.
                                                                                                                                                                                                                                                                                                                                                                          (GEHO-) GEN
                                                                                                                                                                                                                                                                                                                                                                                                                                          14-APR-1986;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22-OCT-1987.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    monoclonal antibody - used for prepn. of thrombolytic drug increased thrombolytic activity and specificity and dreactivity to fibrinogen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GHRPLDKKR 9
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                                                                                                                                                                                                                                                                                                                                                                             HOSPITAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (updated)
(first entry)
                                                                                                                                                                                                                                                                                                                                 Haber E;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     monoclonal antibody;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
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                                                                                                                                                      English.
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Pred.
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hes 0;
                                                                                                           fibrinogen cross-reactivity
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                                                                                                                                                                                                                      screening
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. Monoclonal
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They have increased binding to in vitro and in vivo MAbs can be used in immunoassays for fibrin in the fibrinogen or other proteins. They can be used as i ligands for the purification of fibrin.

(Updated on 03-OCT-2002 to add missing OS field.)

vivo thrombi.

presence of

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RESULT 7
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                                                                                  Query Match
Best Local Similarity
Matches 7; Conserv
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Best Local
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                                                                                                                                                                       Sequence
                                                                                                                                                                                                          Peptide is epitopic to fibrin, allowing anti-fibrin Abs raised without cross-reactivity to fibrinogen. They are particularly useful in detection of fibrin and thrombi.
                                                                                                                                                                                                                                                                                                                                       Fibrin-specific monoclonal antibodies - lacking fibrinogen cross-reactivity, obtd. using peptide(s) comprising fibrin-specific epitopic sequences.
                                                                                                                                                                                                                                                                                               Claim 1; Page 17; 12pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1990-185723/24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matsueda
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23-APR-1984;
30-JAN-1986;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-JAN-1986;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Peptide antigenic to fibrin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         fibrin; fibrinogen; thrombi; immunoaffinity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAR05558 standard; peptide; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                        ш
                                        GHRPLDK 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GHRPLDK 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GHRPLDK 7
                                                                                                                                                                       7 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        7
                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          80.4%; So llarity 100.0%; I Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Haber E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          84US-0603155.
86US-0824228.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            86US-0824228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /label=Lys or Arg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /label=Asp or Glu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   label=Leu or Val
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  'label=Pro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        'label=His or Pro
                                                                         80.4%; Score 41; DB 11; L
100.0%; Pred. No. 7.8e+05;
... Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                or Val
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 41; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 8;
5. 7.8e+05;
0;
                                                                                                                   Length 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 7;
                                                                           Indels
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                                                                      Gaps
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                                                                         0,
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RESULT 9 AAR65792

AAR65792 standard; peptide; 7 AA

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1 GHRPLDK 7

Conservative

Mismatches

Indels

0;

Gaps

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RESULT 8
AAR63269
ID AAR6
XX AAR6
XX AAR6
XX Inhi
KW hiru
KW COOO
KW haem
OS Synt
XX IO-N
PF 03-M
XX OYA
PF 03-M
XX OYA
PF 03-M
XX OYA
PF 10-N
                                                                                                                                                                                                                This sequence represents beta-peptide which was used to raise the monoclonal antibody 59D8. The antibody binds fibrin and may be used in the chimeric molecule of the invention. The chimeric molecule further comprises a thrombin inhibitor linked to the fibrin-binding antibody through a covalent linkage. The chimeric molecule allows fibrin-specific antibody targetting of hirudin and other thrombin thibitors, which is more potent than thrombin on its own. The epitope to which 59D8 binds becomes available only after thrombin cleaves fibrinopeptide B. The chimeric protein may be used for preventing coagulation of the blood. Anti-thrombin targetting can be esp. useful in highly thrombogenic situations such as coronary stent implantation and can be used as an adjunctive therapy with highly selective thrombolytic agents. The thrombin inhibitor is localised to sites of thrombolytic agents. The thrombin inhibitor is localised to sites of thrombin activity by the antibody which binds to thrombin but does inhibition allows the total amount of thrombin inhibitor used to be substantially reduced, resulting in a reduced potential for generalised
             Query Match
Best Local S
Matches 7
                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Fibrin-binding antibody linked to thrombin inhibitor - useful for preventing blood coagulation by specifically targetting inhibitor to site of thrombin activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; Page 13; 53pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1994-358195/44.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bode C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        03-MAY-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       03-MAY-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO9425491-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Inhibitor; thrombin; chimeric molecule; fibrin-binding; antibody; hirudin; monoclonal; 59D8; fibrinopeptide B; coagulation; P79; coronary stent implantation; adjunctive therapy; fibrinogen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Beta-peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (HARD ) HARVARD COLLEGE. (UYEM-) UNIV EMORY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-NOV-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    haemorrhage.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21-JUL-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAR63269;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAR63269 standard; peptide; 7
             Local Similarity les 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Haber E, Runge M;
                                                                                                                                          7 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   94WO-US04881
100.0%; ++
                                      80.4%; Score 41;
100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   raise monoclonal antibody 59D8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A
      ; DB 15; I
D. 7.8e+05;
cches 0;
                                                                     Length 7;
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RESULT 10
AAP82686
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local S
Matches 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               epitopic sequences, they can be used to prepare hybridoma cell lines, which produce antifibrin-specific monoclonal antibodies substantially devoid of fibrinogen cross-reactivity. These antibodies are useful in the in vivo and in vitro detection of thrombi and fibrin deposits.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic epitopic peptide(s) of variable length - capable of eliciting fibrin specific antibodies free of fibrinogen cross-reactivity.
                                               08-OCT-1986;
                                                                                                                                                                                                                                                                                                         05-DEC-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              riprin-specific epitopic peptides; thrombi antifibrin-specific monoclonal antibodies
    08-OCT-1986;
                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                    Fibrin;
                                                                                                                                                                                                                                                               Human fibrin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 2; Column 18; 12pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1994-332411/41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            23-APR-1984;
30-JAN-1986;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAR65792;
                                                                                                                                    JP63093800-A
                                                                                                                                                                                                                                                                                                                                                    AAP82686
                                                                                                                                                                                                                                                                                                                                                                                          AAP82686 standard; peptide; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Haber
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24-AUG-1992;
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                                                                                         25-APR-1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GEHO)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26-JUN-1995
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                                                                                                                                                                                                                      beta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GHRPLDK 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GHRPLDK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A
A
                                                                                                                                                                                                                                                            beta chain N-terminal
                                                                                                                                                                                                                    chain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                       (first
                                               86JP-0237876
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  89US-0454954.
92US-0932729.
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86US-0824228.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    epitopic peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matsueda
                                                                                                                                                                                                                  N-terminal; monoclonal antibody; thrombosis
                                                                                                                                                                                                                                                                                                       entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          80.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score
Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          41;
No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 15; I
7.8e+05;
hes 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       fibrinogen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
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Query Match
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Best Local
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    prepd. f
    and mouse

                                                                Human fibrin beta-chain peptides A and B were synthesised and coupled to BSA for injection into mice. The peptides were used to raise antibodies to human fibrin. Monoclonal antibodies specific for fibrin are used in the production of bispecific monoclonal antibodies which also recognise truncated tPA muteins lacking the finger, EGF and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 anti-fibrin; monoclonal
antithrombotic agent; bi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tissue plasminogen activator; t-PA; anti-fibrin; monoclonal antibody; hy antithrombotic agent; bispecific and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The
Sequence
                                             Kringle
                                                                                                                                                                                                     Example 1; Page 14; 38pp; Japanese.
                                                                                                                                                                                                                                                                            having
                                                                                                                                                                                                                                                                                                 Hybrid
                                                                                                                                                                                                                                                                                                                                                 WPI; 1993-408334/51
                                                                                                                                                                                                                                                                                                                                                                                                                                    20-JUN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-JUN-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              JP05304992-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human fibrin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAR44837;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; Page 1; 8pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New anti-human fibrin monoconal antibody for diagnosing thrombosis - prepd. from hybridoma obtd. by applying cell fusion mouse spleen and mouse myeloma cells, and measures human fibrin.
                                                                                                                                                                                                                                                       reduced
                                                                                                                                                                                                                                                                                                                                                                                            (TAKE ) TAKEDA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19-NOV-1993.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-JUN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (MITN ) MITSUBISHI GAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          peptide is linked to a carrier protein via a linking agent d to immunise mice. The mouse spleen cells and amyeloma ce can be fused to produce a hybridoma secreting MAbs which be can be fused to produce the control of the control of
                                                                                                                                                                                                                                               monoclonal antibody - used for prepn. of increased thrombolytic activity and spec: d reactivity to fibrinogen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               μ
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                                               domains.
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  AA;
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A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                beta-chain N-terminal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first
                                                                                                                                                                                                                                                                                                                                                                                            CHEM
                                                                                                                                                                                                                                                                                                                                                                                                                                      91JP-0148936
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 92JP-0158301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     peptide;
                                                                                                                                                                                                                                                                                                                                                                                              IND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               80.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CHEM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 41;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      줒
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; mutein; fibrin; antigen; hybridoma; thrombolysis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 9;
3. 7.8e+05;
0;
                                                                                                                                                                                                                                                                       n. of thrombolytic specificity and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
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                                                                                                                                                                                                                                                                                                 drug
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Similarity 7; Conserv

Conservative

80.4%; but 100.0%; Pr

Score 41; DB; Pred. No. 7.8

.8e+05; 14;

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Gaps

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Length 8; Indels

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AAR63270
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                                                                          Query Match
Best Local
                                                          Matches
                                                                                                                                                                                                coagulation of the blood. Anti-thrombin targetting the control of the blood. Anti-thrombin targetting the control of the blood in highly thrombogenic situations such as coronary stent implantation and can be used as an adjunctive therapy with highly selective thrombolytic agents. The thrombin inhibitor is localised to sites thrombolytic agents. The thrombin inhibitor is localised to sites
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  preventing b
                                                                                                                                                     of thrombin activity by the antibody which binds to thrombin but do not cross react with uncleaved fibrinogen. The selectivity of inhibition allows the total amount of thrombin inhibitor used to be substantially reduced, resulting in a reduced potential for general:
                                                                                                                                                                                                                                                                                      This sequence represents a synthetic beta-peptide which was used to immunopurify the monoclonal antibody 59D8 which was raised against beta-peptide (see also ARA63269). The antibody binds fibrin and may be used in the chimeric molecule of the invention. The chimeric molecule further comprises a thrombin inhibitor linked to the fibrin-binding antibody through a covalent linkage. The chimeric molecule allows fibrin-specific antibody targetting of hirudin and other thrombin inhibitors, which is more potent than thrombin on its own. The epitope to which 59D8 binds becomes available only after thrombin cleaves
                                                                                                                    Sequence
                                                                                                                                               haemorrhaging.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Fibrin-binding antibody linked to thrombin inhibitor - useful for preventing blood coagulation by specifically targetting inhibitor to site of thrombin activity
                                                                                                                                                                                                                                                                                    fibrinopeptide B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1994-358195/44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bode C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   03-MAY-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 03-MAY-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Inhibitor; thrombin; chimeric molecule; fibrin-binding; antibody; hirudin; monoclonal; 59D8; fibrinopeptide B; coagulation; P79; coronary stent implantation; adjunctive therapy; fibrinogen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (HARD ) HARVARD COLLEGE. (UYEM-) UNIV EMORY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10-NOV-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO9425491-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     haemorrhage.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic beta-peptide used to raise monoclonal antibody 59D8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAR63270;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAR63270 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21-JUL-1995
                                                                       Local
                           ш
                                                         Similarity 7; Conser
GHRPLDK
                           GHRPLDK 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Haber E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GHRPLDK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GHRPLDK
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Page 14;
                                                         Conservative
                                                                                                                  ΑĄ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 93US-0058699
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              94WO-US04881
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                                                                                                                                                                                                                                                             The chimeric protein may be used for preventing blood. Anti-thrombin targetting can be esp. useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Runge
                                                                      80.4%;
100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  53pp; English.
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                                                      0;
                                                                 Score 41;
Pred. No.
                                                      Mismatches
                                                  DB 15; I
7.8e+05;
thes 0;
                                                                             Length 8;
                                                  Indels
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AAR65793
ID AAR6
XX AAR6
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                          RESULT 14
AAR44838
ID AAR44
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Best Local S
Matches 7
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     20-JUN-1994
                                                     AAR44838;
                                                                                             AAR44838 standard; peptide; 12 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ARA65789-R65794 are synthetic peptides comprising fibrin-specific epitopic sequences, they can be used to prepare hybridoma cell lines, which produce antifibrin-specific monoclonal antibodies substantially devoid of fibrinogen cross-reactivity. These antibodies are useful in the in vivo and in vitro detection of thrombi and fibrin deposits.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        23-APR-1984;
30-JAN-1986;
22-DEC-1989;
24-AUG-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic epitopic eliciting fibrin sp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAR65789-R65794 are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 3; Column 18; 12pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cross-reactivity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1994-332411/41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Haber E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     23-APR-1984;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Fibrin-specific epitopic peptides; thrombi detection; fibrinogen; antifibrin-specific monoclonal antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18-OCT-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US5357042-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAR65793;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAR65793 standard; peptide; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Fibrin-specific epitopic peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-JUN-1995
                                                                                                                                                                                                                                                                                                                                Similarity 7; Conserv
                                                                                                                                                                                                                                                                   GHRPLDK 7
                                                                                                                                                                                                                                GHRPLDK 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hui K,
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                                                                                                                                                                                                                                                                                                                                   Conservative
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                   A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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86US-0824228.
89US-0454954.
92US-0932729.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   84US-0603155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ic peptide(s) of variable length - capa specific antibodies free of fibrinogen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note= "May be absent may be bonded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matsueda
                                                                                                                                                                                                                                                                                                                      100.0%; +-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GR
                                                                                                                                                                                                                                                                                                                        Score 41; DB 15; 1; Pred. No. 7.8e+05; 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      and if present
to keyhole limpet
                                                                                                                                                                                                                                                                                                                                                                         Length 8;
                                                                                                                                                                                                                                                                                                                           Indels
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Human fibrin beta-chain peptide

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                                                                                                                                                                                                                                                                                                                                     RESULT 15
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Best Local S
Matches 7
          WPI; 2000-566116/53
                                                                                                                                                                                                                                                                  19-JAN-2001
                                                                                                                                                                                                                                                                                                            AAB10666 standard; peptide; 11 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human fibrin beta-chain peptides A and B were synthesised and coupled to BSA for injection into mice. The peptides were used to raise antibodies to human fibrin. Monoclonal antibodies specific for fibrin are used in the production of bispecific monoclonal antibodies which also recognise truncated tPA muteins lacking the finger, EGF and which also recognise truncated tPA muteins lacking the finger,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hybrid monoclonal antibody - used for prepn. of thrombolytic drug having increased thrombolytic activity and specificity and reduced reactivity to fibrinogen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-NOV-1993.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          JP05304992-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tissue plasminogen activator; t-PA; mutein; fibrin; antigen; anti-fibrin; monoclonal antibody; hybridoma; thrombolysis antithrombotic agent; bispecific antibody.
                                                                                                                                     17-AUG-2000
                                                                                                                                                                                  Unidentified
                                                                                                                                                                                                                                                                                        AAB10666;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1993-408334/51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20-JUN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                               Lahann J, Lendlein A;
                                                                                        12-FEB-1999;
                                                                                                                                                            DE19905793-A1
                                                                                                                                                                                                        bridging molecule
                                                                                                                                                                                                                   Fibrinogen-alpha; polymer-encapsulated ingredient; drug release;
                                                                                                                                                                                                                                          Fibrinogen-beta peptide fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Kringle 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; Page 14; 38pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17-JUN-1992;
                                                      (LAHA/) LAHANN J.
(LEND/) LENDLEIN A.
                                                                                                              12-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (TAKE ) TAKEDA CHEM IND LTD
                                                                                                                                                                                                                                                                                                                                                                                             μ
                                                                                                                                                                                                                                                                                                                                                                       9
                                                                                                                                                                                                                                                                                                                                                                                                                   7;
                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                            GHRPLDK 7
                                                                                                                                                                                                                                                                                                                                                                       GHRPLDK 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      domains
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                (first entry)
                                                                                          99DE-1005793
                                                                                                               99DE-1005793
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      91JP-0148936
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            92JP-0158301
                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                             Score 41; DB 14; Length 12; Pred. No. 0.085;
                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                   0,
                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                  0;
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peptide, nucleic acid, polysaccharide, glycosaminoglycan, terpene or lipid bridging molecules into a water-swellable polymer (e.g. an acrylic polymer or polyvinylpyrrolidone), optionally protecting the bridging molecules against reaction with the cleaving reagent, by reaction with a carbonate, chloroformate, azidoformate or dicarbonate ester, thermally or photochemically activating the bridging molecules to crosslink the polymer chains, and introducing the crosslinked material into the body so that the bridging molecules will be cleaved by the reagent (e.g. an enzyme) at the intended release site, optionally after changing the ph to deprotect the bridging molecules or contacting the material with an enzyme that renders the bridging molecules cleavable by the cleaving managers are accepted by the cleaving molecules or contacting the material with an enzyme that renders the bridging molecules cleavable by the cleaving
                                                     reagent, especially a racemase, epimerase or exopeptidase. This sequences a fragment of fibrinogen-alpha which acts as a chemically active bridge which is used in the method of the invention.
Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              bridging molecules into the polymer during its manufacture, where the bridging molecules are cleavable by interaction with an endogenous reagent at the intended release site. The method is especially useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This invention describes a novel method for the selective release of polymer-encapsulated ingredients comprising incorporating reactive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 3; 6pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                bridging molecules
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Selective release of polymer-encapsulated ingredients, especially drugs, comprises crosslinking the polymer chains with cleavable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    for drug release in vivo, preferably by
11 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The method is especially useful by incorporating amino acid,
                                                                                                                       This sequence
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Search completed: June 27, 2003, 17:44:04 Job time: 70 secs Query Match
Best Local Similarity
Matches 6; Conserv თ GHRPLD 6 Conservative 70.6%; bu 100.0%; Pre Score 36; Pred. No. Mismatches DB 21; . 0.84; 0 Indels 0 Gaps

Length 11

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Result
No.
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Maximum
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Maximum Match
Listing first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Database
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Perfect score:
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BG
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seq length:
  protein search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                 Published Applications AA:*

1: /cgn2_6/ptodata/1/pubpaa/US08 NEW PUB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/PCT_NEW PUB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/US06 NEW PUB.pep:*

4: /cgn2_6/ptodata/1/pubpaa/US07 NEW PUB.pep:*

4: /cgn2_6/ptodata/1/pubpaa/US07 PUBCOMB.pep:*

6: /cgn2_6/ptodata/1/pubpaa/US07 PUBCOMB.pep:*

6: /cgn2_6/ptodata/1/pubpaa/US07 PUBCOMB.pep:*

7: /cgn2_6/ptodata/1/pubpaa/US09 PUBCOMB.pep:*

9: /cgn2_6/ptodata/1/pubpaa/US09 PUBCOMB.pep:*

9: /cgn2_6/ptodata/1/pubpaa/US09 PUBCOMB.pep:*

10: /cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep:*

11: /cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep:*

12: /cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep:*

13: /cgn2_6/ptodata/1/pubpaa/US00_NEW_PUB.pep:*

14: /cgn2_6/ptodata/1/pubpaa/US60_NEW_PUB.pep:*
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Gapop 10.0 , Gapext 0.
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51
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  GHRPLDKKR 9
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Copyright (c) 1993 - 2003 Compugen Ltd
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             US-10-142-935-9
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US-10-054-569-5
US-09-954-385-130
US-09-982-877-278
US-09-983-802-349
US-10-083-357-1073
US-09-981-877-383
US-10-225-567A-2064
US-09-881-572A-3
US-10-040-862-10000
US-10-040-862-10018
US-10-040-862-10018
US-10-040-862-10018
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        Sequence 9, Appli
Sequence 8, Appli
Sequence 5, Appli
Sequence 130, App
Sequence 278, App
Sequence 290, App
Sequence 349, App
Sequence 3173, App
Sequence 383, App
Sequence 383, Appli
Sequence 373, Appli
Sequence 9737, Appli
Sequence 9737, Appli
Sequence 10018, A
Sequence 1018, A
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Sequence 27,
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22	22	22	22	22	23	23	23	23	23	23	24	24	24	24	24	24	24	24	24	24	24	24	24	24	24
43.1	43.1	43.1	43.1	43.1	45.1	45.1	45.1	45.1	45.1	45.1	47.1	47.1	47.1	47.1	47.1	47.1	47.1	47.1	47.1	47.1	47.1	47.1	47.1	47.1	47.1
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US-09-947-137-21	US-09-954-166-15	US-09-935-682-29	US-10-097-065-642	US-10-318-780-12	US-10-223-711-2	US-09-821-687-5	US-10-225-567A-920	US-09-876-904A-247	.US-09-876-904A-478	US-09-821-687-7	US-10-211-088-250	US-08-424-550B-593	US-09-880-748-2853	US-10-100-861-4	US-09-880-748-2538	US-09-880-748-2513	US-09-880-748-2480	US-09-880-748-2444	US-09-880-748-2369	US-10-185-815-20	US-09-984-057-56	US-09-984-056-56	US-09-192-854-36	US-09-782-672-49	US-09-968-561A-54
21,	Sequence 15, Appl	e 29,	Sequence 642, App	Sequence 12, Appl	Sequence 2, Appli	e 5,	Sequence 920, App	Sequence 247, App	478,	Sequence 7, Appli	Sequence 250, App	Sequence 593, App	28	4,	Sequence 2538, Ap	Sequence 2513, Ap	Sequence 2480, Ap	Sequence 2444, Ap	Sequence 2369, Ap	Sequence 20, Appl	Sequence 56, Appl	Sequence 56, Appl	Sequence 36, Appl	Sequence 49, Appl	Sequence 54, Appl

ALIGNMENTS

RESULT 1 US-10-142-935-9

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NUMBER OF SEQ ID NOS: 11
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 9
; LENGTH: 10
; TYPE: PRT
; ORGANISM: Staphylococcus epidermidis US-10-142-935-9
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US-10-142-935-8
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           Sequence 8, Application US/10142935

Publication No. US20030044418A1

GENERAL INFORMATION:
APPLICANT: DAVIS, Stacey
APPLICANT: HOOK, Magnus A.O.
TITLE OF INVENTION: METHOD AND COMPOSITIONS FOR INHIBITING THROMBIN-INDUCED COAGULAT
FILE REFERENCE: P07201US01/BAS
CURRENT APPLICATION NUMBER: US/10/142,935
CURRENT FILING DATE: 2002-05-13
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Best Local S
Matches 6
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Publication No. US20030044418A1

GENERAL INFORMATION:

APPLICANT: DAVIS, Stacey

APPLICANT: HOOK, Magnus A.O.

TITLE OF INVENTION: METHOD AND COMPOSITIONS FOR INHIBITING THROMBIN-INDUCED COAGULAT

FILE REFERENCE: P07201US01/BAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/10/142,935
CURRENT FILING DATE: 2002-05-13
PRIOR APPLICATION NUMBER: US 60/290,072
PRIOR FILING DATE: 2001-05-13
PRIOR APPLICATION NUMBER: US 60/290,072
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100.0%; Pr
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                    Publication No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
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                                                                                       FILE REFERENCE: GC690
CURRENT APPLICATION NUMBER: US/09/954,385
CURRENT FILING DATE: 2001-09-12
NUMBER OF SEQ ID NOS: 433
SOFTWARE: FastSEQ for Windows Version 4.0
EQ ID NO 130
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OTHER INFORMATION: binding peptide
                                                                                                                                                                                                                                                                                                                                 APPLICANT: Aehle, Wolfgang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/10/068,569
CURRENT FILING DATE: 2002-02-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR FILING DATE: 2001-05-13
NUMBER OF SEQ ID NOS: 11
SOFTWARE: PatentIn version 3.1
                                    ORGANISM: Artificial Sequence
                                                             LENGTH: 12
TYPE: PRT
                                                                                                                                                                                                                                                   PPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILE REFERENCE: 480140.475
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Srinivasula, Srinivasa M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT
ORGANISM: Mus musculus
                                                                                                                                                                                             PPLICANT: Winetzky, Deborah S.
ITLE OF INVENTION: Binding Phenol Oxidizing Enzyme-peptide
ITLE OF INVENTION: Complexes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 15
TYPE: PRT
ORGANISM: Staphylococcus epidermidis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PPLICANT: Alnemri, Emad S. (TLE OF INVENTION: A CONSER (TLE OF INVENTION: CASPASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
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                                                                                                                                                                                                                                                                                                                                                    INFORMATION:
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                                                                                                                                                                                                                                                            Janssen, Giselle G.
Murray, Christopher J.
                                                                                                                                                                                                                                                                                Van Gastel, Franciscus J.C.
Janssen, Giselle G.
                                                                                                                                                                                                                                                                                                                 Baldwin, Toby
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Pred. No.
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RESULT 6
US-09-948-783-290
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Best Local Similarity
Whiches 4; Conserve
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LENGTH: 15
TYPE: PRT
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 09/892,877
PRIOR FILING DATE: 2001-06-28
PRIOR APPLICATION NUMBER: 09/437,658
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILE REFERENCE: PZ028P1
CURRENT APPLICATION NUMBER:
CURRENT FILING DATE: 2001-0
                                                                                                                                                                                                                                                                                                                     PRIOR FILING DATE: 2000-09-
                                                                                                                                                                                                                                                                                                                               CURRENT FILING DATE: 2001-09-10 PRIOR APPLICATION NUMBER: 60/231,846
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PRIOR FILING DATE: EARLIER FILING DATE: 1999-11-10
NUMBER OF SEQ ID NOS: 461
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                TLE OF INVENTION: 97 Human secreted proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PPLICANT: Ruben et. al.
                                                                FILING DATE: 1998-01
APPLICATION NUMBER:
APPLICATION NUMBER: 60/085,924
                 APPLICATION NUMBER: 60/085,906 FILING DATE: 1998-05-18
                                                  FILING DATE: 1998-05-18
                                                                                                APPLICATION NUMBER: 60/085,180
                                                                                                                 FILING DATE: 1998-05-12
                                                                                                                                                               APPLICATION NUMBER: 60/085,094
                                                                                                                                                                                    APPLICATION NUMBER: 60/085,093 FILING DATE: 1998-05-12
                                                                                                                                                                                                                               FILING DATE: 1999-11-10
APPLICATION NUMBER: PCT/US99/09847
                                                                                                                               APPLICATION NUMBER: 60/085,105
                                                                                                                                                                                                                   LING DATE: 1999-05-06
                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/09/948,783
                                                                                                                                                  LING DATE:
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5. US20030077809A1
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Pred. No.
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99;
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77;
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0

NUMBER: 60/085,922

EARLIER

DATE: 1997-07-08

60/055,722

EARLI

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PRIOR FILING DATE: 1998-05-18
PRIOR APPLICATION NUMBER: 60/085,921
PRIOR FILING DATE: 1998-05-18
PRIOR FILING DATE: 1998-05-18
PRIOR PRIOR PRIOR DATE: 1998-05-18
PRIOR PRIOR DATE: 1998-05-18
PRIOR FILING DATE: 1998-05-18
PRIOR PRIOR DATE: 1998-05-18
PRIOR PRIOR DATE: 1998-05-18
PRIOR FILING DATE: 1998-05-18
PRIOR FILING DATE: 1998-05-18
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; ORGANISM: Homo sapiens US-09-948-783-290 SEQ ID NO 290 NUMBER OF SEQ ID NOS: 465 TYPE: PRT CENGTH: PatentIn Ver. 2.0

밁 Query Match Best Local Matches 9 GHRP 12 4; Similarity Conservative 51.0%; Score 26; 100.0%; Pred. No. 0 Mismatches DB 99; 9: Length 15; Indels 0

Gaps

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US-09-983-802-349 Sequence 349, Application US/09983802 Publication No. US20030022185A1 TITLE OF INVENTION: INFORMATION: Fischer et al 123 Human Secreted Proteins

APPLICATION NUMBER: DATE: 1997-07-08 APPLICATION NUMBER: APPLICATION NUMBER: APPLICATION NUMBER: 1998-07-07 1999-01-08 PCT/US98/13684 09/227,357 60/051,926

CURRENT APPLICATION NUMBER: US/09/983,802
CURRENT FILING DATE: 2001-10-25
PRIOR APPLICATION NUMBER: EARLIER APPLICATION
PRIOR FILING DATE: EARLIER FILING DATE: 1999
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PRIOR APPLICATION NUMBER: EARLIER APPLICATION
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PR APPLICATION NUMBER: DATE: 1997-07-08 APPLICATION NUMBER: APPLICATION NUMBER: 1997-07-08 1997-07-08 1997-07-08 60/052,803 60/051,929 60/052,793 60/051,925

APPLICATION NUMBER: APPLICATION NUMBER: DATE: 1997-07-08 1997-07-08 60/051,916 60/051,931 60/052,732 60/051,932

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> ; ORGANISM: Homo sapiens US-09-983-802-349 SEQ ID NO 349 Query Match Best Local PRIOR PRIOR PRIOR SOFTWARE: PRIOR PRIOR PRIOR PRIOR TYPE: PRT ENGTH: FILING DATE: FILING DATE: EARLIER APPLICATION NUMBER: E Ξ APPLICATION NUMBER: FILING DATE: EARLIE FILING DATE: APPLICATION NUMBER: EARLIER FILING DATE: EARLIER FILING FILING DATE: FILING DATE: EARLIER APPLICATION NUMBER: APPLICATION NUMBER: EARLIER FILING DATE: FILING DATE: FILING DATE: EARLIER FILING APPLICATION NUMBER: EARLIER APPLICATION NUMBER: EARLIER APPLICATION NUMBER: EARLIER FILING DATE: EARLIER FILING FILING DATE: APPLICATION NUMBER: APPLICATION NUMBER: FILING DATE: EARLIEF APPLICATION NUMBER: EARLIER FILING DATE: EARLIER FILING APPLICATION NUMBER: EARLIER APPLICATION APPLICATION NUMBER: EARLIER APPLICATION NUMBER: EARLIER FILING DATE: EARLIEF APPLICATION NUMBER: LING DATE: SEQ ID NOS: 672 PatentIn NUMBER: EARLIER EARLIER FILING NUMBER: EARLIER EARLIER FILING EARLIER 51.0%; 83.3%; EARLIER EARLIER EARLIER Score 26; I Pred. No. 1 APPLICATION NUMBER: DATE: 1997-09-12 APPLICATION NUMBER: DATE: 1997-08-18 APPLICATION NUMBER: DATE: 1997-08-18 DATE: 1997-09-12 APPLICATION NUMBER: APPLICATION NUMBER: DATE: 1997-08-18 APPLICATION NUMBER: APPLICATION NUMBER: APPLICATION NUMBER: APPLICATION NUMBER: DATE: 1997-08-18 APPLICATION NUMBER: APPLICATION NUMBER: DATE: 1997-08-18 DATE: 1997-08-18 DATE: 1997-08-18 APPLICATION NUMBER: DATE: 1997-08-18 APPLICATION NUMBER: DATE: 1997-08-18 DATE: 1997-08-18 APPLICATION NUMBER: DATE: 1997-08-18 APPLICATION NUMBER: APPLICATION NUMBER: 1997-09-12 Length 60/058,661 60/058,785 60/058,660 60/058,664 60/055,954 60/055,984 60/055,684 60/056,360 60/055,964 60/055,947 60/055,950 60/055,953 60/055,949 60/055,948

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Similarity

RESULT 8 US-10-083-357-1073 GENERAL INFORMATION: Sequence 1073, Application US/10083357 Publication No. US20030054370A1

APPLICANT: Qiandong Zeng et al.
TITLE OF INVENTION: Systemic Discovery of New Genes
FILE REFERENCE: 032796-090

CURRENT APPLICATION NUMBER: US/10/083,357
CURRENT FILING DATE: 2002-02-27
NUMBER OF SEQ ID NOS: 1346

SEQ ID NO 1073

US-10-083-357-1073 LENGTH: 18
TYPE: PRT
ORGANISM: Saccharomyces cerevisiae

Query Match
Best Local S Matches Similarity 51.0%; 5c. 100.0%; Pr Score 26; pred. No. Mismatches DB 9; .2e+02; Length 18; °.

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PRIOR APPLICATION NUMBER: 09/437,658
PRIOR PRILING DATE: 1999-11-10
PRIOR APPLICATION NUMBER: PCT/US99/09847
PRIOR FILING DATE: 1999-05-06
PRIOR FILING DATE: 1999-05-06
PRIOR APPLICATION NUMBER: 60/085,093
PRIOR FILING DATE: 1998-05-12
PRIOR APPLICATION NUMBER: 60/085,094
PRIOR FILING DATE: 1998-05-12
                         PRIOR APPLICATION NUMBER: 60/08
PRIOR FILING DATE: 1998-05-18
PRIOR APPLICATION NUMBER: 60/08
PRIOR FILING DATE: 1998-05-18
PRIOR APPLICATION NUMBER: 60/08
PRIOR APPLICATION NUMBER: 60/08
PRIOR FILING DATE: 1998-05-18
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; ORGANISM: Homo sapiens
US-09-892-877-383
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                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 60/
PRIOR FILING DATE: 1998-05-12
PRIOR APPLICATION NUMBER: 60/
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CURRENT FILING DATE: 2001-09-10
PRIOR APPLICATION NUMBER: 60/231,846
PRIOR FILING DATE: 2000-09-11
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PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US/09/437,658
PRIOR FILING DATE: EARLIER FILING DATE: 1999-11-10
NUMBER OF SEQ ID NOS: 461
SOFTWARE: Patentin Ver. 2.0
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TITLE OF INVENTION: 97 Human secreted proteins
FILE REFERENCE: PEZO28P1
CURRENT APPLICATION NUMBER: US/09/892,877
CURRENT FILING DATE: 2001-06-28
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  APPLICATION NUMBER:
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o. US20030100051A1
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60/085,925
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Pred. No. 1.3e+02;
2; Mismatches 2
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APPLICANT: Roth, Monica;
APPLICANT: Bupp, Keith;
APPLICANT: University of Medicine and Dentistry of New Jersey
TITLE OF INVENTION: Targeting Viral Vectors to Specific Cells
FILE REFERENCE: 601-1-095
CURRENT APPLICATION NUMBER: US/09/881,572A
CURRENT FILING DATE: 2002-03-05
PRIOR APPLICATION NUMBER: 60/212,239
PRIOR APPLICATION DATE: 50070-06-16
NUMBER OF SEQ ID NOS: 27
SOFTWARE: PatentIn version 3.0
SEQ ID NO 3
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US-09-881-572A-3
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Best Local Similarity
                                                                                                                                                                                                                                                                                              Sequence 3, Application US/09881572A Patent No. US20020164583A1
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APPLICANT:
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SEQ ID NO 383
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Best Local
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PRIOR FILING DATE: 2000-12-19
NUMBER OF SEQ ID NOS: 2292
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ORGANISM: moloney murine leukemia virus
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TITLE OF INVENTION: ANTIGENIC PEPTIDES AND ANTIBODIES FOR G PROTEIN-COUPLED RECEPTOR
                          TYPE:
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ORGANISM: Homo sapiens
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PRIOR APPLICATION NUMBER: 60/0
PRIOR FILING DATE: 1998-05-18
NUMBER OF SEQ ID NOS: 465
                                        ENGTH:
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ORGANISM: Homo sapiens
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100.0%; Pred. NO. 1.3e+(
tive 0; Mismatches •
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Pred. No. 1
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US-10-4-1

US-10-662-9737

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US-10-040-862-10000
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US-10-040-862-9737
Sequence 10000, Application US/10040862 Publication No. US20030078396A1 GENERAL INFORMATION:
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Best Local
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Best Local
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CURRENT FILING DATE: 2001-11-06
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PRIOR FILING DATE: 2000-03-01
PRIOR APPLICATION NUMBER: US 60/190,479
PRIOR FILING DATE: 2000-03-17
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                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US 60/223,378 FILING DATE: 2000-08-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US 60/222,903
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CATION NUMBER: US 60/223,416
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Retter, Marc
Corixa Corporation
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NUMBER: US 60/218,950
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Pred. No. 3.
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Pred. No. 3
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Matches
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SEQ ID NO 10000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                        APPLICANT: Mannion, Jane
APPLICANT: Retter, Marc
APPLICANT: Coriva Corporation
TITLE OF INVENTION: Compositions and Methods for the Detection, Diagnosis and Therap
TITLE OF INVENTION: Hematological Malignancies
FILE REFERENCE: 014058-013520US
                                                                                                                                                                                                            APPLICANT: Gaiger, Alexander
APPLICANT: Algate, Paul A.
APPLICANT: Mannion, Jane
APPLICANT: Retter, Marc
APPLICANT: Corixa Corporati
                 CURRENT APPLICATION NUMBER: US/10/040,862 CURRENT FILING DATE: 2001-11-06 PRIOR APPLICATION NUMBER: US 60/186,126 PRIOR FILING DATE: 2000-03-01 PRIOR APPLICATION NUMBER: US 60/190,479 PRIOR FILING DATE: 2000-03-17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IITLE OF INVENTION: Compositions and Methods for the Detection, Diagnosis and Therap
IITLE OF INVENTION: Hematological Malignancies
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
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APPLICATION NUMBER: US 60/200,545
FILING DATE: 2000-04-27
APPLICATION NUMBER: US 60/200,545
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 2000-04-28
APPLICATION NUMBER: US 60/200,999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US 60/200,303 FILING DATE: 2000-04-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US 60/190,479
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US 09/796,692 FILING DATE: 2001-03-01
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APPLICATION NUMBER: US 60/222,903
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                                                                                                                                                                                                                                                                                                                          INFORMATION:
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2000-08-07
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Pred. No. 3.8e+05;
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PRIOR FILING DATE: 2000-04-27
PRIOR APPLICATION NUMBER: US 60/200,303
PRIOR FILING DATE: 2000-04-28
PRIOR APPLICATION NUMBER: US 60/200,799
PRIOR APPLICATION NUMBER: US 60/200,999
PRIOR APPLICATION NUMBER: US 60/200,099
PRIOR APPLICATION NUMBER: US 60/200,099
PRIOR APPLICATION NUMBER: US 60/200,099
PRIOR APPLICATION NUMBER: US 60/200,004
PRIOR APPLICATION NUMBER: US 60/200,201
PRIOR APPLICATION NUMBER: US 60/218,950
PRIOR PILING DATE: 2000-05-02
PRIOR APPLICATION NUMBER: US 60/222,903
PRIOR FILING DATE: 2000-06-03
PRIOR PILING DATE: 2000-08-03
PRIOR PILING DATE: 2000-08-03
PRIOR PILING DATE: 2000-08-03
PRIOR APPLICATION NUMBER: US 60/223,416
PRIOR APPLICATION NUMBER: US 60/223,378
PRIOR PILING DATE: 2000-08-07
PRIOR APPLICATION NUMBER: US 60/223,378
PRIOR PILING DATE: 2000-08-07
PRIOR APPLICATION NUMBER: US 60/223,378
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PRIOR APPLICATION NUMBER: US 60/223,378
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PRIOR APPLICATION NUMBER: US 60/223,378
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Result
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Maximum Match 100%
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DB seq length: 19
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/cgn2_6/ptodata/1/iaa/backfiles1.pep:*
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   GenCore version 5.1.6 (c) 1993 - 2003 Compugen Ltd
 summaries
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Sequence 10, Appl
Sequence 34, App
Sequence 234, App
Sequence 234, App
Sequence 234, App
Sequence 21, Appl
Sequence 27, Appl
Sequence 27, Appl
Sequence 27, Appl
Sequence 27, Appl
Sequence 122, App
Sequence 8, Appli
Sequence 9, Appli
Sequence 9, Appli
Sequence 9, Appli
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Sequence 10, Appl
Sequence 11, Appl
Sequence 30, Appl
Sequence 79, Appl
Sequence 41, Appl
Sequence 9, Appli
Sequence 9, Appli
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US-08-448-547-1
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Sequence 32,	Sequence 3, A	Sequence 3, A	Sequence 10,	Sequence 3, A	Sequence 10,	417	Sequence 9, A	Sequence 9, A;	Sequence 39,	Sequence 593,	Sequence 306,	Sequence 306,	Sequence 17,	Sequence 17,	8	Sequence 8, A	Patent No. 539576
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ALIGNMENTS

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STRANDEDNESS:
TOPOLOGY: lin
MOLECULE TYPE:
US-08-448-547-1
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Ver

CURRENT APPLICATION NUMBER: US/08/448,547

FILING DATE: 30-JUN-1995

CLASSIFICATION: 435

PRIOR APPLICATION DATA:

APPLICATION NUMBER: PCT/JP94/01844

FILING DATE: 01-NOV-1994

PRIOR APPLICATION DATA:

APPLICATION NUMBER: PCT/JP94/01844

PRIOR APPLICATION DATA:

APPLICATION NUMBER: JP 5-297325

FILING DATE: 02-NOV-1993

ATTORNEY/AGENT INFORMATION:

NAME: NAKAMUTA, Dean H.

BETSCTENEY NUMBER: DEAN H.

BETSCTE
                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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CITY: Washington
STATE: D.C.
COUNTRY: USA
ZIP: 20037
                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Nakamura, Dean H.
REGISTRATION NUMBER: 33,981
REFERENCE/DOCKET NUMBER: Q-
TELECOMMUNICATION INFORMATION:
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TITLE OF INVENTION: ANTI-HUMAN SOLUBLE FIBRIN ANTIBODY,
TITLE OF INVENTION: HYBRIDOMA, AND IMMUNOASSAYING METHOD
                                                                                                                                                                      LENGTH: 10 amino TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE:
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Inuzuka, Kimiko
Ito, Yumiko
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(202)293-7860
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                                                                                           GENERAL INFORMATION:
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APPLICANT: Edgar I
APPLICANT: Christo
APPLICANT: Marsch
                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
APPLICANT: Edga. .... APPLICANT: Christoph Bode
APPLICANT: Christoph Bode
APPLICANT: Marschall S. Runge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: WordPerfect (Version 5.1)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/058,699
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
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PPLICANT: Marschall S. Runge
ITLE OF INVENTION: FIBRIN-TARGETED INHIBITORS
ITLE OF INVENTION: THROMBIN
                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (617) 542-8906
                                                                                                                                                                                                                                                                                                                                             STRANDEDNESS:
TOPOLOGY: Li
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REGISTRATION NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 05433/004001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: N/A FILING DATE: N/A
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CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PS/2 Model 50Z or 55
OPERATING SYSTEM: MS-DOS (Version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                             LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE:
                                                                                                            No.
                                                                                                                                                                                                                                                                  Similarity 7; Conserv
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                                                                                                                                                                                                                                                                                                                                                                              AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Janis K. Fraser,
                                                                                                                                                                                                                                   GHRPLDK 7
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                                                                                                                         Application US/08058699
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      225 Franklin Street
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                                                                                                                                                                                                                                                                                                                                               Linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19930503
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Pred. No. 0.00062;
); Mismatches 0;
                                                                                                                                                                                                                                                                    Mismatches
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                                                                                                                                                                                                                                                                                           Length 7;
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                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 30,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                    COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                        CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,
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                                                                                   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                        SOFTWARE:
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US-07-876-941A-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 19930503
CLASSIFICATION: 530
RAIOR APPLICATION DATA:
APPLICATION NUMBER: N/A
FILING DATE: N/A
ATTORNEY/AGENT INFORMATION:
NAME: Janis K. Fraser, Ph.D.
REGISTRATION NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 05433/004001
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTEX.
ZIP: 02.110-2804
ZIP: 02.110-2804
COMPUTER READABLE FORM:
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 ru-
MEDIUM TYPE: M.S-DOS (Version 5.0)

Version 5.1)
                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEX: 200154
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
                                                                                       CORRESPONDENCE ADDRESS:
                                                                                                                                   TITLE OF INVENTION:
                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                APPLICANT:
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                                                      ADDRESSEE: LO
COUNTRY: U
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CITY: Boston
                                                                                                                       TLE OF INVENTION:
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                             Palo Alto
: CA
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               USA
                                                                                                                                                Mitchell,
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                                                                                                                                                                Bradley, Danı
Tam, Albert W
                                                                                                                                                                                               Reyes, Gregory R.
                                                                                                                                                                                                                                                                                                                                                                                Conservative
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                                                                      Dehlinger & Associates
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                                                          Cambridge Avenue, Suite 250
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                                                                                                                                                                                                                                                                                                                                                                       80.4%; UT
100.0%; Pr/
0;
                                                                                                                  Antibodies
                                                                                                                                                                                 Daniel W.
                                                                                                                                   Hepatitis E Virus Peptide Antigen and
                                                                                                                                                Carl
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Pred. No. 1.9e+(
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PatentIn Release #1.0, Version #1.25

01-MAY-1992

US/07/876,941A

PRIOR APPLICATION DATA: PRIOR APPLICATION DATA
APPLICATION NUMBER:

FILING DATE:

17-JAN-1992

US 822,335

CLASSIFICATION:

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                                                                                                                                                                                                                                                                                                                   Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 46
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 324-0880
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US 20:
FILING DATE: 17-JUNE-1988
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 11-APRI PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 505,888 FILING DATE: 05-APRIL-1990 PRIOR APPLICATION DATA:
                        SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                    TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
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APPLICANT: YEN, R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ANTI-SENSE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local
                                                        MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                STREET: Two Embarcac
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US 420,9:
FILING DATE: 13-OCTOBER-1989
                                                                                                                            COUNTRY: UZIP: 94111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: si
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Sholtz, Charles REGISTRATION NUMBER: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER:
APPLICATION NUMBER: US/08/747,137 FILING DATE: 12-NOV-1996
                                                                                                                                                                                    ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, 8th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ENGTH:
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5. 5945033
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(415) 324-0960
TO NO: 30:
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                                                                                                                                                                                                                                      ON: NON-CROSSLINKED PROTEIN PARTICLES FOR ON: THERAPEUTIC AND DIAGNOSTIC USE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 2; Length 18;
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US-08-556-597-84
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Best Local Similarity
                                                                                                                               APPLICATION NUMBER: US/08/556,597
FILING DATE:
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/406,330
FILING DATE: 17-MAR-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO:
                TELEFAX: (716) 263-16
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                       COMPUTER: IBM PC Compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOTTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Miller, Jonathan L.
APPLICANT: Lyle, Vicki A.
TITLE OF INVENTION: MIMOTOPES AND ANTI-MIMOTOPES OF
TITLE OF INVENTION: HUMAN PLATELET GLYCOPROTEIN Ib/IX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 4 amino acids
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APPLICATION NUMBER: US 0
FILING DATE: 15-JAN-1991
                                                              TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-576-0200
FORMATION FOR SEQ ID NO: 79:
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SEQUENCE CHARACTERISTICS:
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                                                                                REFERENCE/DOCKET NUMBER: 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Apple, Randolph T
REGISTRATION NUMBER: 36,
REFERENCE/DOCKET NUMBER:
                                              TELEPHONE:
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FILING DATE: 14-MAR-1994
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: New York
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                                                                                                                  Timian, Susan J
                                                                                                                                                                                                                                                                                                                                                                                                                              Clinton Square, P.O. Box 1051
                                                                                                                                                                                                                                                                                                                                                                              USA
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                                                (716)
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                              263-1600
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100.0%; Pr
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                                                                                             34,103
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                                                                                  20884/101
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US-08-592-500-41
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; MOLECULE TYPE:
US-08-556-597-84
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APPLICANT: Lanza,
                                                                                                                                                                                                                                                                                            TELEFAX: (415) 326-24 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                    MOLECULE TYPE: peptide HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 536
PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/592,500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 09-JUL-1993
ATTORNEY/AGENT INFORMATION:
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APPLICANT: Phillips, David R.
APPLICANT: Cazenave, Jean-Pierre
TITLE OF INVENTION: Platelet Glycoprotein V Gene and Uses
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                                                                                                                        LOCATION: 1..11
OTHER INFORMATION:
OTHER INFORMATION:
                                 OTHER INFORMATION: OTHER INFORMATION:
                                                                                                                                                                                                                               LENGTH: 11 amino acids
TYPE: amino acid
TOPOLOGY: unknown
                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: 29
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                                                                                                           OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
      VAME/KEY:
                                                                               NAME/KEY:
                                                                                                                                                                  NAME/KEY: Peptide
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71.4%;
                                                                                                        /note= "Amino acid
human fibrinogen (F
cleavage site."
                               /note= "Amino acid residues identical to GPV."
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Pred. No.
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RESULT 8
US-08-195-006-41
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Best Local
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SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
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                                                                                                                                 FEATURE:
                                                                                                                                                                                                                   HYPOTHETICAL: NO FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/089,455
FILING DATE: 09-JUL-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                           LOCATION: 1..11
OTHER INFORMATION:
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                              MOLECULE TYPE:
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APPLICANT: Cazenave, Jean-Pierre
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OTHER INFORMATION: OTHER INFORMATION:
                                                                      OTHER INFORMATION: OTHER INFORMATION:
                                                                                                               NAME/KEY:
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                                                                                                                                                                                                                                                                                                                                                                                          NAME: Dow, Karen B. REGISTRATION NUMBER:
                              LOCATION:
                                          NAME/KEY:
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                                        Region
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SYSTEM: PC-DOS/MS-DOS
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10-FEB-1994
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100.0%; Pr
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                                                                                                                                          /note= "Amino acid sequence of the
human fibrinogen (Fg) B-beta chain
cleavage site."
/note= "Amino acid residue
identical to GPV."
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identical to GPV."
                                                                     /note= "Amino acid residues
identical to GPV."
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10; Mismatches
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46;
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                                                      Query Match
Best Local S
Matches 4
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION: TELEPHONE: (415) 326-2400
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MEDIUM TYPE: Floppy
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NUMBER OF SEQUENCES:
                                                                                                                                                                                FEATURE:
                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: peptide HYPOTHETICAL: NO
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OTHER INFORMATION:
OTHER INFORMATION:
                                                                 Local Similarity
                                                                                                                         OTHER INFORMATION: OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Dow, Karen B.
REGISTRATION NUMBER: 29,684
REFERENCE/DOCKET NUMBER: 01
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LOCATION:
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OPERATING SYSTEM: PC-DOS/MS-DOS
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GHRP 11
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                                                   51.0%; Score 26; DB 5;
100.0%; Pred. No. 46;
tive 0; Mismatches
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                                                                                                                         /note= "Amino acid residue
identical to GPV."
                                                                                                                                                                                                                                                                 /note= "Amino acid sequence of the human fibrinogen (Fg) B-beta chain cleavage site."
                                                                                                                                                                                            /note= "Amino acid residues
identical to GPV."
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Pred. No.
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GENERAL INFORMATION:
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       APPLICANT: Hagstrom, James E.
APPLICANT: Sebestyen, Magdoina G.
APPLICANT: Wolff, Jon A.
APPLICANT: Wolff, Jon A.
TITLE OF INVENTION: A METHOD FOR COVALENT ATTACHMENT OF COMPOUNDS TO GENES
FILE REFERENCE: Covalent Attachment to Genes
CURRENT APPLICATION NUMBER: US/08/990,015A
CURRENT FILING DATE: 1997-12-12
EARLIER APPLICATION NUMBER: 60/050842
EARLIER APPLICATION NUMBER: 60/050842
EARLIER FILING DATE: 1997-05-26
NUMBER OF SEQ ID NOS: 11
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                               APPLICANT: Budker, Vladimir
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APPLICATION NUMBER: US 08/092,200
FILING DATE: 14-JUL-1993
ATTORNEY/AGENT INFORMATION:
NAME: KOENIG, NATHAN P.
REGISTRATION NUMBER: 38,210
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: 13054.00110
TELECOMMUNICATION INFORMATION:
TELEPHONE: (510) 763-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                 y Match 51.0%;
Local Similarity 44.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: FILING DATE: 07-JUN CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WordPerfect 5.1/ASCII
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                                                                                                                                                                                                             INFORMATION:
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PatentIn
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1999 HARRISON ST
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Ver.
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Pred. No. 55;
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PCT-US94-07916-10

FCT-US94-07916-10

; Sequence 10, Application PC/TUS9407916

; GENERAL-INFORMATION:

APPLICANT: SZOKA, FRANCIS C.
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                                                                   RESULT 13
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                                                                                                                                                         Matches
                                                                                                                                                                     Query Match
Best Local :
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: KOENIG, NATHAN P.
REGISTRATION UNDER: 38,210
REFERENCE/DOCKET NUMBER: 1305
TELECHONE: 1510) 763-2000
TELEPHONE: 1510) 763-2000
TELEFAX: (510) 273-8832
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                                                                                                                                                                                                           TYPE: amino acids
STRANDEDNESS: single
TOPOLOGY: lin-
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ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: WordPerfect 6.1/ASCII CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy
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                                                                                                                                                                   Local
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                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: FILING DATE: 06-JUN CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: Windows 3.11
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                                                                                                                                                                   Similarity
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6113946
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                                                                                                                                                       Conservative
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99 HARRISON ST
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Pred. No. 55;
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Pred. No.
                                                                                                                                                       Mismatches
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PCT-US94-07916-10
                                                                 Matches
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NAME: BERLINER, ROBEK:

NAME: BERLINER, ROBEK:

REGISTRATION NUMBER: 20,121

REFERENCE/DOCKET NUMBER: 5555

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION 1000

TELECOMMUNICATION 2000

TELECOMMUNICATION 2000

TELECOMMUNICATION 2000

TELECOMMUNICATION 2000

TELECOMMUNICATION 2000

TELECOMMUNICATION 2000
                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: HENSLER, ITITLE OF INVENTION: TITLE OF INVENTION: TITLE OF INVENTION: NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: WordPerfect 5.1 / ASCII CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                     TYPE: amino ac
STRANDEDNESS:
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CITY: I
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                                                                                                                                                                                            ENGTH:
2 GYGPKDKRK 10
                              1 GHRPLDKKR 9
                                                                             Similarity
                                                                                                                                                                           amino acid
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201 NORTH FIGUEROA STREET
                                                               Conservative
                                                                                                                                                linear
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SELF-ASSEMBLING POLYNUCLEOTIDE
DELIVERY SYSTEM COMPRISING DENDRIMER
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                                                                             Score 26;
Pred. No.
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                                                             Mismatches
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                                                                                         Length 13;
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US-09-227-357-349 GENERAL INFORMATION: Sequence 349, App. Patent No. 634258 EARLIER APPLICATION NUMBER: 60/052,803
SARLIER FILING DATE: 1997-07-08
SARLIER APPLICATION NUMBER: 60/052,803 CURRENT APPLICATION NUMBER: US/09/227,357 CURRENT FILING DATE: 1999-01-08 APPLICANT: Fischer et al. TITLE OF INVENTION: 123 FILE REFERENCE: PZ010P1 EARLIER APPLICATION NUMBER: 60/051,929 EARLIER FILING DATE: 1997-07-08 RARLIER FILING DATE: SARLIER FILING DATE: 1997-07-08
SARLIER APPLICATION NUMBER: 60/051,925 EARLIER APPLICATION NUMBER: 60/051,926 EARLIER FILING DATE: 1997-07-08 EARLIER APPLICATION NUMBER: PCT/US98/13684 EARLIER FILING DATE: 1998-07-07 PARLIER APPLICATION NUMBER: 60/052,793 APPLICATION NUMBER: 60/052,732 FILING DATE: 1997-07-08 APPLICATION NUMBER: 60/051,931 FILING DATE: 1997-07-08 Application US/09227357 123 Human Secreted Proteins 1997-07-08 1999-01-08

60/051,932

APPLICATION NUMBER: 60/051,916

1997-07-08

EARLIER

APPLICATION NUMBER: 60/051,918 FILING DATE: 1997-07-08 APPLICATION NUMBER: 60/051,930

1997-07-08

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RESULT 15
US-08-827-962-2
; Sequence 2, Application US/08827962A
; Patent No. 6258944
; GENERAL INFORMATION:
; APPLICANT: MERCK & CO., INC.
TITLE OF INVENTION: OB RECEPTOR ISOFORMS AND NUCLEIC ACIDS
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-227-357-349
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Best Local Similarity
Matches 5; Conserv
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SEQ ID NO 349
LENGTH: 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 60/058,664
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APPLICATION NUMBER: 60/055,949
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FILING DATE: 1997-07-08
APPLICATION NUMBER: 60/051,928
FILING DATE: 1997-07-08
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                                                                                                                                                                                                                                     4 PLDKKR 9
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PLICATION NUMBER: 60/055,950
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83.3%;
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FILE REFERENCE: 19693

CURRENT APPLICATION NUMBER: US/08/827,962A

CURRENT FILING DATE: 1997-05-06

PRIOR APPLICATION NUMBER: 60/016,899

PRIOR FILING DATE: 1996-05-06

NUMBER OF SEQ ID NOS: 21

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 2

LENGTH: 19

TYPE: PRT

ORGANISM: Rattus No. 6258944vegicus

US-08-827-962-2
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                                                              Query Match
Best Local Similarity
Matches 5; Conserv
                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION:
10 GHKDLISKR 18
                             1 GHRPLDKKR 9
                                                              Conservative
                                                                                                                                                                                                                                                                                                                      ENCODING THEM
                                                                             51.0%;

    Mismatches

                                                                             Score 26; DB
Pred. No. 81;
                                                                                          DB 4;
                                                                                          Length 19;
                                                            Indels
                                                            0;
```

Gaps

Search completed: June 27, 2003, 17:47:17 Job time: 27 secs

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